

SEQUENCE LISTING

<110> LEVINE, et al.

<120> VARIANTS Of PROTEIN KINASES

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<150> 09/724,676

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<151> 2000-06-15

<150> 135619

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<170> PatentIn version 3.0

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cggaagatgg aggcaaatac ccagttatat ttaagcatgg agatgattta cgtcaagatc	1980
aacttattct tcaaatcatt tcaactcatgg acaagctggt acggaaaagaa aatctggact	2040
tgaaattgac accttataag gtggttagcca ccagtacaaa acatggcctt atgcagttta	2100
tccagtcagt tcctgtggct gaagttcttg atacagaggg aagcattcag aactttttta	2160
gaaaatatgc accaagtgag aatggggccaa atgggattag tgctgaggtc atggacactt	2220
acgttaaaag ctgtgctgga tattgcgtga tcacctatat acttgagatt ggagacaggc	2280
acctggataa ccttttgcta acaaaaacag gaggctgagg caggagaatc gcttgaaccc	2340
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<210> 11
 <211> 1894
 <212> DNA
 <213> Homo sapiens

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gcttgggaagg gaagagagaa caaaagagtt ataaagctgt cctggaagac ccaatgttga	180
agttctcagg actatatcaa gagacatgct ctgatcttta tggtacttgt caagtttttg	240

cagaagggaa gcctttggcc ttgccagtga gaacatccta caaagcattt agtacaagat	300
ggaactggaa tgaatggctg aaactaccag taaaataccc tgacctgccc aggaatgccc	360
aagtggccct caccatatgg gatgtgtatg gtcccggaaa agcagtgctt gtaggaggaa	420
caacggtttc gctctttgga aaatacggca tgtttcgcca agggatgcat gacttgaaag	480
tctggcctaa tgtagaagca gatggatcag aaccacaaa aactcctggc agaacaagta	540
gcaactcttc agaagatcag atgagccgtc ttgccaagct caccaaagct catcgacaag	600
gacacatggg gaaagtagat tggctggata gattgacatt tagagaaata gaaatgataa	660
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atcaagataa agccttgacc aaaatcctga catctgttat ttgggatcta cctcaggggg	1080
ccaaacaggc cttggcactt ctgggggaaat ggaacccgat ggatgtagag gactccttgg	1140
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gacaggccga tgatgaggat ttgttgatgt acctattaca attggtccag gctctcaa	1260
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cagtgtcaga aaatgtgtca aattctggaa taaattctgc agaaatagat agctcccaa	1380
ttataaccag ccccttccct tcagtctctt cacctcctcc tgcataaaaa acaaaagaag	1440
ttccagatgg cgaaaatctg gaacaagatc tctgtacctt cttgatatcg agagcctgca	1500
aaaactcaac actggcta	1560
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cttattaccc ttaagaatta ttttttctca tgctataaac acaaagatga tttaaaataa	1620
gatatgagca atcatcaaaa gtctagatat agtatttata tctaactgta ttccctgaca	1680
gctgggttca tggaaatgct tggtggcatc aacatttctc tccctttcac taaggatata	1740
actttccagg agctaggata taactgggga gggtaatggg tactttcagt ttaatatctg	1800
tttggcatgt aaagtagttg agtatgatat ttcacagttc ttcaggagta tatattttca	1860
cattattttac taaaacattg gaatcattta aaaa	1894

<210> 12
 <211> 750
 <212> DNA
 <213> Homo sapiens

 <220>
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 <222> (1)..(750)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 12
 cagtgnctc cgggccgcg gccgcagcca gcacccgcg cgccgcagct ccgggaccgg 60
 ccccgccgc cgccgccgc atgggcaacg ccgccgcgc caagaagggc agcgagcagg 120
 agagcgtgaa agaattctta gccaaagcca aagaagattt tcttaaaaaa tgggaaagtc 180
 ccgctcagaa cacagccac ttggatcagt ttgaacgaat caagaccctc ggcacgggct 240
 ccttcggggc ggtgatgctg gtgaaacaca aggagaccgg gaaccactat gccatgaaga 300
 tctcgacaa acagaaggcg gtgaaactga aacagatcga acacaccctg aatgaaaagc 360
 gcacctgca agctgtcaac tttcgttcc tcgtcaaact cgagttctcc ttcaaggaca 420
 actcaaactt atacatggtc atggagtacg tgcccgccg ggagatgttc tcacacctac 480
 ggcgatcgg aagggtcagg taagcgggcc acccccatc acatcaggct gtcagggtgt 540
 ccacaggtgg cagtgcacga ccaagcccc tgggaatgca gaggagtcca gcatacttca 600
 acatgcagg ggtctccag accctgtggg tttctgtttc ccctctgctg aggaatatgt 660
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 aagttgctgc caggcgcggt gggtctcacc 750

<210> 13
 <211> 794
 <212> DNA
 <213> Homo sapiens

<400> 13
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 ttctttctcg accctacat tgccctcaat gtggacgact cgcgcatcgg ccaaacggcc 180
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 ggacgcaaga tcgagctggc tgtctttcac gatgccccca taggctacga cgacttcgtg 300
 gccaaactg ccatccagtt tgaggagctg ctgcagaacg ggagccgcca cttcgaggac 360

tggattgatc tggagccaga aggaagagtg tatgtgatca tcgatctctc agggtcgtcg	420
ggtgaagtga aaatacctaa ctctgcattc tgtgaaaggg agagagttga aatgaggcac	480
agctgaagtt tcaccactcc attattctgc catcaagcat cccttcagct cccatctctc	540
aaatgctgca gccccatcaa tggaaaaaaa tacattcatg ggaattgcac aaagtttaca	600
acacatgcaa attccaagga gctcagagag aggtgtgggg gtgagatggt gtgactacac	660
ccccttcttc taggatgcgc cgaaggattc tttttcttgc tcatatgctg ttagaaagcc	720
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cactgcttgc aaaa	794

<210> 14
 <211> 2083
 <212> DNA
 <213> Homo sapiens

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ctctgttttc ttctttccaa aactgtgcac ccctggatga aacctccatc aaggagacc	180
tacaagttgc ctggggttca gtgctctaga aagttccaag gtttgtggct tgaattattc	240
taaagaagct gaaataattg aagagaagca gaggccagct gtttttgagg atcctgctcc	300
acagagaatg ctctgcaccc gttgatactc cagttccaac accatcttct gagatgatcc	360
tgattcccag aatgctcttg gtgctgttcc tgctgctgcc tatcttgagt tctgcaaaag	420
ctcagggttaa tccagctata tgccgctatc ctctgggcat gtcaggaggc cagattccag	480
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tggactcaga agaaggggat ggagcctggt gccctgagat tccagtggaa cctgatgacc	600
tgaaggagtt tctgcagatt gacttgcaca ccctccattt tatcactctg gtggggaccc	660
aggggcgcca tgcaggaggt catggcatcg agtttgcccc catgtacaag atcaattaca	720
gtcgggatgg cactcgctgg atctcttggc ggaaccgtca tgggaaacag gtgctggatg	780
gaaatagtaa cccctatgac attttcctaa aggacttggg gccgcccatt gtagccagat	840
ttgtccggtt cattccagtc accgaccact ccatgaatgt gtgtatgaga gtggagcttt	900
acggctgtgt ctggctagat ggcttgggtg cttacaatgc tccagctggg cagcagtttg	960
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acacgcctgc	tcccagttca	ttgcatctat	ttttagtctc	tgctaacctc	ctgagaagtc	1980
cacattctct	ttctagattt	ggctctgccg	tgctcctggc	ctaatttgag	caactctcct	2040
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 <212> DNA
 <213> Homo sapiens

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tgcaagtgga atgaattatt tgcacctcca taaaattatt catcgtgatc tcaaatacacc	1140
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<210> 16
 <211> 1327
 <212> DNA
 <213> Homo sapiens

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cactgtaatg gttcgtttta atttttcatt ttggattttt ttttaatttc ctttgctact	180
tagattagaa agaacattga tctttcaaac atagatctga atatgaaaga gaaaagaaac	240
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ttggcattcc tttttcacac tttaaaatca ggtaataagt cagtccatct gaggaaagcc	360
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actttgttga acatccttaa gaatgctgat ctcagcttgg cttttcaaac atacacaata	600
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gacagtgttg atggaggtca cgattctgtc attttggatc cagagcgact tgagcctggg	720
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accacagacc tttatagcag ccogtatgat attattcaca ttatgatatg tgtttattat	1260
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gaaaaaa	1327

<210> 17
 <211> 364
 <212> DNA
 <213> Homo sapiens

<400> 17	
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ccctgatccc tcagcgttca tgcagcctct tgtccacgga ggctgggtgcc ctgcatgtgc	180
tgtctgcccgc tcggggcccc gggccccccc agcgcctatc tttctccttt ggggtccctg	240
tccgacctgt gggcgccaat ggcctcccc tgacctcagg gttccttggg ggctgggctg	300
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aaac	364

<210> 18
 <211> 923
 <212> DNA
 <213> Homo sapiens

<400> 18	
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ggggtccagg ctctgctgtg tgcgtgcacc agctgcctcc aggccaacta cacgtgtgag 180
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acctcagggg ctggctcagg taccaagttc ttcagggcat catgtctgtg gttggctttc 660
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agataagata atgggttcac agaaaatctc tcattccgtg ggattcccaa ggtctagtag 840
acaattatcc actacagaag tcctgcctct ctaagctttg caactgtacg tacatctatt 900
aaaattcagc tgggtcccccc tga 923

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<210> 19

<211> 1739

<212> DNA

<213> Homo sapiens

<220>

<221> -

<222> (1)..(1739)

<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 19

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atctgggtggc cctccagatg aaccgacgtc accggatgcc tggatatgag accatgaaga 180
acaaagacac aggtcactca aataggcaga gtgacgtcag aatcaagttc gagcacaacg 240
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actctgaggt gtccagacag gtgcggatca aggcttccca gtccgcaggg gatataaata	540
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aatgctctac taagagattg atggggtgct ggggtggagg gggggaagcc tgnagcccaa	1620
gagaccctgt tcctggnaga atgaatgggg aatattcata aataatgtac acaaagtaac	1680
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<210> 20

<211> 1832

<212> DNA

<213> Homo sapiens

<220>

<221> -

<222> (1)..(1832)

<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 20

ccgccgcccc ggccccggc atgcagcccc ggctgcggag gtgacactca cggaccttag	60
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atctacatta	catgaacaat	gagctctcca	tcctgtgtaa	aaaccaagat	gatcttgata	480
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cccaagtgcc ccccatcaaa cctgggcgcc cg

1832

<210> 21
<211> 1269
<212> DNA
<213> Homo sapiens

<400> 21
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gtggagtgcc ctttttgtga tgaagtttcc aaatacgaga agctcgccaa gatcggccaa 180
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aagatccttc agcttctaaa acacgagaat gtggtcaact tgattgagat ttgtcgaacc 360
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ycccccaaa 1269

<210> 22
<211> 623

<212> DNA
 <213> Homo sapiens

 <220>
 <221> -
 <222> (1)..(623)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 22
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 gggggcctct ctagcttgcg gcctgtgtct atggtcgggc cctctgcgtc cagctgctcc 120
 ggaccgagct cgggtgtatg gggccgtagg aaccggctcc ggggccccga taacgggccg 180
 cccccacagc accccgggct ggcgtgaggg tctcccttga tctgagaatg gctacctctc 240
 gatatgagcc agtggctgaa attggtgtcg gtgcctatgg gacagtgtac aaggcccgtg 300
 atccccacag tggccacttt gtgccctcaa gagtgtgaga gtccccaatg gaggaggagg 360
 tggaggaggc cttcccatca gcacagttcg tgagggtggct ttactgaggc gactggaggc 420
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 ggagatcaag gtaaccctgg tgtttgagca tgtagaccag gacctaagga catatctgga 540
 caaggcaccc ccaccaggct tgccagccga aacgatcaag gtgagtgggg ttggtaggca 600
 ttganagggtg gattgggacc ttt 623

<210> 23
 <211> 502
 <212> DNA
 <213> Homo sapiens

<400> 23
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 ggaccgagct cgggtgtatg gggccgtagg aaccggctcc ggggccccga taacgggccg 180
 cccccacagc accccgggct ggcgtgaggg tctcccttga tctgagaatg gctacctctc 240
 gatatgagcc agtggctgaa attggtgtcg gtgcctatgg gacagtgtac aaggcccgtg 300
 atccccacag tggccacttt gtgccctcaa gagtgtgaga gtccccaccc acctctcctt 360
 ttgaggcttc tccttctcct tccatttct ctacactaag gggatatgttc cctcttgtcc 420
 ctttccctac ctttatatattt ggggtccttt tttatacagg aaaaacaaaa caaagaaata 480
 aagtcgacgc ggccgcgaat tc 502

<210> 24
 <211> 1148
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(1148)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 24
 catcatatga ggtggacagg aagcccagcc aagccatggg ggccgacccc tcccntggat 60
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 cctgggaaga ctccctttctg ctccccaaaa cccaaggcc tggctcgggg ccactggagc 180
 cgcaggcggg acatatgtgt gaccggccct ctgcccctgg cagccccgcg ctgtgtactg 240
 taaggacgtg ctggacatcg agcagttctc cactgtgaag ggcgtcaatc tggaccacac 300
 agacgacgac ttctactcca agttctccac gggctctgtg tccatcccat ggcaaaacga 360
 gatgatagaa acagaatgct ttaaggagct gaacgtgttt ggacctaatg gtaccctccc 420
 gccagatctg aacagaaacc accctccgga accgcccag aaagggtgtc tccagagact 480
 cttcaagcgg cagcatcaga acaattccaa gagttcgcgc agctccaaga ccagttttaa 540
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 ggagcccctg ctctggtggg gctgccaggg gagaccccgg gagccgggga aggaggccgt 720
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 gactcaagac ttccagagcc tcaaatgaga aaatgtcttt attaaatgta gaaagtgatc 1140
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<210> 25
 <211> 1679
 <212> DNA

<213> Homo sapiens

<220>

<221> -

<222> (1)..(1679)

<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 25

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ggcattttcc tgtgtcctcc cacacctgca ctctcccaag gctcttgagg tctctctgag	180
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gcagcatcag aacaattcca agagtccgcc cagctccaag accagtttta accaccacat	1080
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ccccggccgg ggtggattgg atttgtcttt ggtgaacatt gcaatagaaa tccaattgga	1380
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tttttaaaga aaagttttgt aaattttctct actgtctcag tttacatttt gtatatattgt	1500
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gtgttccaag gcatttttagc ggggaggggg ttatcaaaaa aaaaaaaatg tgactcaaga	1620
cttccagagc ctcaaagag aaaatgtctt tattaaatgt agaaagtgat ccatacttc	1679

<210> 26
 <211> 897
 <212> DNA
 <213> Homo sapiens

 <220>
 <221> -
 <222> (1)..(897)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 26	
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ggcattttcc tgtgtcctcc cacacctgca ctctcccaag gctcttggtg tcaactctgag	180
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aggactgggg tggtcagcgg agacttccag gagaaagcct ggggtggggca gggacaccca	840
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<210> 27
 <211> 1224
 <212> DNA
 <213> Homo sapiens

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 <213> Homo sapiens

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<212> DNA
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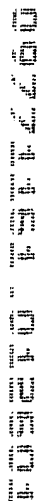
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 ctggaggtga aaatagggga ttttggactg gcaaccaaag tcgaatatga cggggagagg 360
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<212> DNA
<213> Homo sapiens

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<210> 35
<211> 951
<212> DNA
<213> Homo sapiens

<220>
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<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 35

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ggaaggccat ggctctgttg cctccttttc ttgcctctca cagattggaa gtatctaggg      180
acagtgggtg gctaggacag tgctggctgc aggggggtctg ggagcgtkgg cctcacagtg      240
gccttctcta tcctctgcaa caccctccag ccgaattctc aacatacctc aacttctgcc      300
gctccctgcg gtttgacgac aagcccgact actcttacct acgtcagctc ttccgcaacc      360
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agtgaggaga gccccattg gaccagtgtt tgcttagtgt ctctactgta ttttctttaa      900
aaamaaaam aaaaawaaam srcmaaaakw acmacwmaaa aaccagcac a                951

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<210> 36
<211> 2063
<212> DNA
<213> Homo sapiens

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<220>
<221> -
<222> (1)..(2063)
<223> "n" can be any nucleotide 'a', 'c', 'g' or 't'

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<400> 36
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aaccgcccga ggctaccgg gaggacaaga ccgagcctaa ggcctacagg cggcgggcgg      180
ccctcagccc actgggaggc cgggacgaca gcccgtgtc ccacagggcc tctcagagcc      240
tgaggagccg caagtcccc agcccggcag gaggtggcag cagcccctat tctcggcggc      300
tgccgcgctc cccgagcccc tacagtcgcc gccgctcccc cagctacagc cgccacagct      360

```

cctacgagcg gggcgggcgac gtgtccccta gtccctacag cagcagcagc tggcgccgct 420
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attcatctag gcattcaaga tctcgtagca ggcacagatt gtctagatcc agaagtcgtc 540
attctagtat ttctcctagc acactaactc tgaagagtag cctggcagct gaattgaaca 600
agaataaaaa agcacgagca gcagaggcag caagagccgc agaagcagcg aaagctgcag 660
aagcaactaa ggctgctgag gctgctgcca aggctgcaaa agcttcaaac acttctacac 720
ctaccaaggg gaacacggaa actagtgcc gtgcatcaca aacaaaccat gtgaaggatg 780
tgaagaaaat taaaattgaa catgcacctt ctccctcaag tggtggaact ttaaaaaatg 840
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ataaagccac caagaaagca gtcatagtgt gaaaggagag taaatctgct gctacaaagg 960
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agaaggagca acatgtagct ttagtcacct ctacattacc accgttacct ttgcctccca 1080
tgctgcctga agataaagaa gctgatagct tacgaggaaa tatttcagta aaagcagtta 1140
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taccaggagg agatgatctt tcaaagagtc cagaggaaaa gaaaacagca acacagttac 1260
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tgattctggg tcatcagaag ttcataaaac actaaattaa ggtttaaaat aaaaaatgta 1380
cttattcttt caacttaaat atatgaattc tattagagtt ttgaaaaaaa tcataattgc 1440
ttgcattcct attcacaatg actaataaaa ccaaatttaa ttttcwctcg taatttgtaa 1500
tgtattaatt tcctggggta ttgtatatgt taaagatact ttctgttatt aagagattga 1560
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tagcttcaag taagtgccta aagagacctc tttcccttaa aacctgttaa tcagttaaag 1800
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aagactttaa gttagatctg attttagaat tgcagttgag gtagtgcta gtgtgtgaat 1920
ttgaggcat tttctaaact ggccgggcac agtggctcat gcctgtaatc ccagcacttt 1980
gggaggccca ggtgggagaa tcacttgagt ccaggagttt gataccagcc tgagcaacac 2040
aggaacccc atctctacca aaa 2063

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<212> DNA
<213> Homo sapiens

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cggcggcgcc gaggggagga gagcggccca tggaccgcg gggcccggcg cccagactc 180
tgcccgctcg ggacggagcc caagatgtcg gcctagggcg gggcgcgacg acgcgagcgg 240
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<210> 38
<211> 1244
<212> DNA
<213> Homo sapiens

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 <211> 3507
 <212> DNA
 <213> Homo sapiens

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aaagagcttc gagcagtgga agatgtacgg ccacctcaca aagtaacgga ctactcctca	360

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 <211> 3138
 <212> DNA
 <213> Homo sapiens

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<210> 45
<211> 2100
<212> DNA
<213> Homo sapiens

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<220>
<221> -
<222> (1) .. (2100)

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<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 45

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cctcctcctt	ctcagctgct	ctgttctcga	cagcttcggg	gaactgattc	cgcagccttc	180
caatgaagtc	aatctactgg	attcaaaaac	aattcaaggg	gagctgggct	ggatctctta	240
tccatcacat	gggtgggaag	agatcagtg	tgtggatgaa	cattacacac	ccatcaggac	300
ttaccaggtg	tgcaatgtca	tggaccacag	tcaaaaacaat	tggctgagaa	caaactgggt	360
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<210> 46
 <211> 1479
 <212> DNA
 <213> Homo sapiens

 <220>
 <221> -
 <222> (1)..(1479)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 46	
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<210> 48
 <211> 1154
 <212> DNA
 <213> Homo sapiens

<400> 48	
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 tttttttttt aaga 1154

<210> 49
 <211> 930
 <212> DNA
 <213> Homo sapiens

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<210> 50
 <211> 2616
 <212> DNA
 <213> Homo sapiens

<400> 50
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 <213> Homo sapiens

<220>
 <221> -

<222> (1)..(1464)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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<210> 55

<211> 1080
 <212> DNA
 <213> Homo sapiens

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 <223> "n" can be any nucleotide 'a', 'c', 'g' or 't'

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 <211> 1665
 <212> DNA
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<213> Homo sapiens

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<221> -
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<210> 65
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<213> Homo sapiens

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<223> "n" can be any nucleotide 'a', 'c', 'g' or 't'

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<210> 73

<211> 2544

<212> DNA

<213> Homo sapiens

<220>

<221> -

<222> (1)..(2544)

<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 73

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<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(2324)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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 <211> 1396
 <212> DNA
 <213> Homo sapiens

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1396

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<211> 513
<212> DNA
<213> Homo sapiens

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<211> 2044
<212> DNA
<213> Homo sapiens

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 <212> DNA
 <213> Homo sapiens

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 <213> Homo sapiens

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cccaaccagc	ccgtgggagag	tgacgagagc	ctaggcggcc	tctctgctgc	cctgcgctcc	960
tggcacttga	ctccaagctg	ccctctggac	ccagcacccc	tcagggaggc	cggctgtcct	1020
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gtggaaggac	tggcccttgg	cagctctgca	tcatcgtcgt	cagagccacc	gcagattatc	1140
atcaaccctg	cccgcacagaa	gatggtccag	aagctggccc	tgtacgagga	tggggccctg	1200
gacagcctgc	agctgctgtc	gtccagctcc	ctcccaggct	tgggcctgga	acaggacagg	1260
cagggggcccg	aagaaagtga	tgaatttcag	agctgatgtg	ttcacctggg	cagatcccc	1320
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tctgccggca	gtgggggtgg	gggcccattg	ccgcggggga	gagaaggagg	tggccctgct	1440
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catgatgaga	ccctgtctct	gccaaaaaat	tttttaaact	attagcctgg	cgtggtagcg	1920
acgcttgtgg	tcccagctgc	tggggaggct	gaagtaggag	gatcatttat	gcttgggagg	1980

tcgaggctgc agtgagtcac gattgtatga ctgcactcca gcctgggtga cagagcaaga	2040
ccctgtttca aaaagaaaaa ccctgggaaa agtgaagtat ggctgtaagt ctcatgggttc	2100
agtcctagca agaagcgaga attctgagat cctccagaaa gtcgagcagc acccacctcc	2160
aacctcgggc cagtgtcttc aggcctttact ggggacctgc gagctggcct aatgtggtgg	2220
cctgcaagcc aggccatccc tgggcgccac agacgagctc cgagccaggt caggcttcgg	2280
aggccacaag ctgagcctca ggcccaggca ctgattgtgg cagagggggcc actaccaag	2340
gtctagctag gcccagacc tagttacaga cagggaaggg aaacattttc agggaaaaga	2400
catgtatcac atgtcttcag aagcaagtca ggtttcatgt aaccgagtgt cctcttgctg	2460
gtccaaaagt agcccagggc tgtagcacag gcttcacagt gattttgtgt tcagccgtga	2520
gtcacactac atgccccctg gaagctgggc attggtgacg tccaggttgt ccttgagtaa	2580
taaaaacgta tgttgcaatc tcgggaaa	2608

<210> 82
 <211> 1237
 <212> DNA
 <213> Homo sapiens

<400> 82	
cgcggaacccg gccggccccag gccgcgccc gccgcggccc tgagaggccc cggcaggtcc	60
cggccccggcg gcggcagcca tggccggggg gccgggcccg ggggagcccg cagcccccg	120
cgcccagcac ttcttgtagc aggtgccgcc ctgggtcatg tgccgcttct acaaagtgat	180
ggacgccttg gagcccccg actggtgcca gtgcgccgcc ctgatcgtgc gcgaccagac	240
cgagctgcgg ctgtgcgagc gctccgggca gcgcacggcc agcgtcctgt ggccctggat	300
caaccgcaac gcccggtgtg cgcacctcgt gcacatcctc acgcacctgc agctgctccg	360
tgcgcgggac atcatcacag cctggcacc tcccgccccg cttccgtccc caggcaccac	420
tgccccgagg ccagcagca tcctgcacc cgccgaggcc gaggcctgga gcccccgaa	480
gttgccatcc tcagcctcca ccttcctctc ccagctttt ccaggctccc agaccattc	540
agggcctgag ctcggcctgg ttccaagccc tgcttccctg tggcctccac cgccatctcc	600
agcccccttct tctaccaagc caggcccaga gagctcagt tccctcctgc agggagcccc	660
ccccctctcg ttttgctggc ccctctgtga gatttcccg ggcaccaca acttctcgga	720
ggagctcaag atcggggagg gtggctttgg gtgcgtgtac cgggcggtga tgaggaacac	780
ggtgtatgct gtgaagaggc tgaaggagaa cgctgacctg gaggggactg cagtgaagca	840

gagcttcctg accgaggtgg agcagctgtc caggtttcgt cacccaaaca ttgtggactt	900
tgctggctac tgtgctcaga acggcttcta ctgcctgggtg tacggcttcc tgcccaacgg	960
ctccctggag gaccgtctcc actgccagac ccaggcctgc ccacctctct cctggcctca	1020
gcgactggac atccttctgg gtacagcccg agcaagtcag gtttcatgta accgagtgtc	1080
ctcttgctg tccaaaagta gccccagggt gtagcacagg cttcacagtg attttgtggt	1140
cagcgtgag tcacactaca tgccccctg aagctgggca ttggtgacgt ccagggtgtc	1200
cttgagtaat aaaaacgtat gttgcaatct cgggaaa	1237

<210> 83
 <211> 1286
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(1286)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 83	
gaggggacct taaaaattac cggccacaaa aagaaaataa atttaggaaa aattattggt	60
ttatTTTTagg gtgatgttca cagaagaaga tgtcaaattc tacttggtg aacttgact	120
tgctttagac catctacata gcttggaat aatttataga gacttaaaac cagaaaatat	180
acttcttgat gaagaaggtc acatcaagtt aacagatttc ggcctaagta aagagtctat	240
tgaccatgaa aagaaggcat attctTTTTg tggaactgtg gagtatatgg ctccagaagt	300
agttaatcgt cgaggtcata ctacagagtgc tgactgggtg tcttttggtg tgttaatgtt	360
tgaaatgctt actggtacac tccctttcca aggaaaagat cgaaaagaaa caatgactat	420
gattcttaaa gccaaacttg gaatgccaca gtttttgagt cctgaagcgc agagtctttt	480
acgaatgctt ttcaagcgaa atcctgcaaa cagattaggt gcaggaccag atggagttga	540
agaaattaaa agacattcat ttttctcaac gatagactgg aataaactgt atagaagaga	600
aattcatccg ccatttaaac ctgcaacggg caggcctgaa gatacattct attttgatcc	660
tgagtttact gcaaaaactc ccaagattc acctggcatt ccacctagt ctaatgcaca	720
tcagcttttt cgggggttta gttttgttgc tattacctca gatgatgaaa gccaaagctat	780
gcagacagtt ggtgtacatt caattgttca gcagttacac aggaacagta ttcagtttac	840
tgatggatat gaagtaaaag aagatattgg agttggctcc tactctgttt gcaagagatg	900

tatacataaa gctacaaaca tggagtttgc agtgaaggta aatTTTTTTT atttaaaatg	960
caattcatac agttcttggt catgcatgtc agtaccagtt aaaaattaca ctccccctgt	1020
tgttaaaagt gccttttggt ataaaaaagt taaatatctg gctagtgatc ttcagagatc	1080
ttaatctaga accctgtgag ctaaaggtaa ggtgggtata tatctagttt tcccagagca	1140
gtagcagttt acacctcaag tgattttttt tcttttttta cctcaagtga tttttaaagt	1200
atctttttac tctgagaagt ccccatTTTA tgctcanggt gtcagcaaT tCctcaaaat	1260
tgtgtgcaaa attttgtatg ttTaca	1286

<210> 84
 <211> 752
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(752)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 84	
atgccgctgg cgcagctggc ggacccgtgg cagaagatgg ctgtggagag cccgtccgac	60
agcgctgaga atggacagca aattatggat gaacctatgg gagaggagga gattaaccca	120
caaactgaag aagtcagtat caaagaaatt gcaatcacac atcatgtaaa ggaaggacat	180
gaaaaggcag atccttccca gtttgaactt ttaaaagtat tagggcaggg atcatttggga	240
aagggtttct tagttaaaaa aatctcaggc tctgatgcta ggcagcttta tgccatgaag	300
gtattgaaga aggccacact gaaagttcga gaccgagttc ggacaaaaat ggaacgtgat	360
atcttggtag aggttaatca tccttttatt gtcaagttgc attatgcttt tcaaactgaa	420
gggaagttgt atcttatttt ggattttctc aggggaggag atttgtttac acgcttatcc	480
aaagaggtga tgttcacaga agaagatgtc aaattctact tggctgaact tgcacttgct	540
ttagaccatc tacatagcct gggaataatt tatagagact taaaaccaga aaagtaagga	600
atcatgctac taagttgaat acaatgtaat atgattgttt aggagattat aaaaaatcaa	660
gtggcttcat gaaactccca cagtaatgtn tagcgtgcct gtgcttcaca tctctgctaa	720
cactgtagtt tcatacttta aatnactcag tt	752

<210> 85
 <211> 1826
 <212> DNA

<213> Homo sapiens

<400> 85

cgagcgcggc gcccttgagc tgcaccgcgg cgcaggtttg cgagccgact tgtcagccgg	60
ccaagaaaag gaagctccgt cccttcccgc tcaccgcggt tccccacccc ttgtactcta	120
aactctgcas agggcgagcg ygcggccack gakgcgccga ggaggagcga gcgccgccgg	180
gcagcggcgt gccctcgggg gagagggcgc cggakargag cggcggcgcg gcggcgakgg	240
cgcggcgcgc gatggcagct gcttagcccc gcgggcgcgg agcagccccg agctgtggct	300
ggccaggcgg tgcggctggg cgggggacgc cgcgcgcgtt gctgcccggc ccggagagat	360
gagcacggag gcggacgarg gcatcacttt ctctgtgcc aacctcgccc cctcgggctt	420
ctgcaccatc cccgagggcg gcatctgcag gaggggagga gcggcggcgg tgggcgaggg	480
cgaggagcac cagctgccac cgcgcgcgcc gggcagyttc tggaacgtgg agagcgccgc	540
tgcccctggc atcggttgtc cggcggccac ctctcgagc agtgccaccc gaggccgggg	600
cagctctgtt ggcgggggca gccgacggac cacggtggca tatgtgatca acgaagcgag	660
ccaagggcaa ctggtggtgg ccgagagcga ggccctgcag agcttgcggg aggcgtgcga	720
gacagtgggc gccaccctgg aaaccctgca ttttgggaaa ctcgactttg gagaaaccac	780
cgtgctggac cgcttttaca atgcagatat tgcggtggtg gagatgagcg atgccttccg	840
gcagccgtcc ttgttttacc accttggggg gagagaaaagt ttcagcatgg ccaacaacat	900
catcctctac tgygatacta actcggactc tctgcagtca ctgaaggaaa tmatttgcca	960
gaagaatact atgtgcactg ggaactacac ctttgttcct tacatgataa ctccacataa	1020
caaagtctac tgctgtgaca gcagcttcat gaaggggttg acagagctca tgcaaccgaa	1080
cttcgagctg cttcttggac ccatctgctt acctcttggt gatcgtttta ttcaactttt	1140
gaaggtggca caagcaagtt ctagccagta cttccgggaa tctatactca atgacatcag	1200
gaaagctcgt aatttataca ctggtaaaga attggcagct gagttggcaa gaattcggca	1260
gcgagtagat aatatcgaag tcttgacagc agatattgtc ataaatctgt tactttccta	1320
cagagatata caggactatg attctattgt gaagctggta gagactttag aaaaactgcc	1380
aacctttgat ttggcctccc atcaccatgt gaagtttcat tatgcatttg cactgaatag	1440
gagaaatctc cctggtgaca gagcaaaagc tcttgatatt atgattcca tggtgcaaag	1500
cgaaggacaa gttgcttcag atatgtattg cctagtgggt cgaatctaca aagatatgtt	1560
tttgactct aatttcacgg aactgaaaag cagagaccat ggagcttctt ggttcaaaaa	1620

ggcatttgaa tctgagccaa cactacagtc aggaattaat tatgcggtcc tcctcctggc 1680
agctggacac cagtttgaat ctccctttga gctccggaaa gttggtaatt acaacttgat 1740
atttctacat ggaaatcaag aaactcggac ccaacttggt gcaaagacgg atctccgccg 1800
attctgacgg ctctccaggt tttgtc 1826

<210> 86
<211> 476
<212> DNA
<213> Homo sapiens

<400> 86
gccggcgggtg gcgcggcgga gaccgcggtg gtataacaag aggattgcct gatccagcca 60
agatgcagag cacttctaata catctgtggc ttttatctga tatttttaggc caaggagcta 120
ctgcaaattgt ctttcgtgga agacataagt ggatgttcaa atgagagaat ttgaagtgtt 180
gaaaaaactc aatcacaaaa atattgtcaa attatttgct attgaagagg agacaacaac 240
aagacataaa gtacttatta tggaaatttg tccatgtggg agtttataca ctgttttaga 300
agaaccttct aatgcctatg gactaccaga atctgaattc ttaattgttt tgcgagatgt 360
ggcgggtgga atgaatcatc tacgagagaa tggatatagt caccgtgata tcaagccagg 420
aaatatcatg cgtgcactat accattctct cgtagatgat tcattccacc caccac 476

<210> 87
<211> 2131
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(2131)
<223> "n" can be any nucleotide 'a', 'c', 'g' or 't'

<400> 87
gaattattgc tttggaagaa aancttacat ctccacaaga ccttgcttcc caaataagac 60
atgtcagaag atgctgataa tcctcacttc tgcattacaa atcgctcata ggtgcatctg 120
caggatctta cttggatcca gagtgctagc tgcaaaagct tctgggaatt gtacacttaa 180
ctctgaagtt ttatctttta tataggaagt gccgcgtatg atgctgtcct tgacagaaat 240
gtggccatta agaagctcag cagacccttt cagaaccaa cacatgccaa gagagcgtac 300
cgggagctgg tcctcatgaa gtgtgtgaac cataaaaaca ttattagttt attaaatgtc 360
ttcacacccc agaaaacgct ggaggagttc caagatgttt acttagtaat ggaactgatg 420

gatgccaaact tatgtcaagt gattcagatg gaattagacc atgagcgaat gtcttacctg	480
ctgtaccaaaa tgttgtgtgg cattaagcac ctccattctg ctggaattat tcacagggat	540
ttaaaaccaa gtaacattgt agtcaagtct gattgcacat tgaaaatcct ggactttgga	600
ctggccagga cagcaggcac aagcttcatg atgactccat atgtggtgac acgttattac	660
agagcccctg aggtcatcct ggggatgggc tacaaggaga acgtggatat atggtctgtg	720
ggatgcatta tgggagaaat ggttcgccac aaaatcctct ttccaggaag ggactatatt	780
gaccagtggga ataaggtaat tgaacaacta ggaacaccat gtccagaatt catgaagaaa	840
ttgcaacca cagtaagaaa ctatgtggag aatcggccca agtatgcggg actcaccttc	900
cccaaactct tcccagattc cctcttccca gcggactccg agcacaataa actcaaagcc	960
agccaagcca gggacttgtt gtcaaagatg ctagtgattg acccagcaaa aagaatatca	1020
gtggacgacg ccttacagca tccctacatc aacgtctggt atgaccacgc cgaagtggag	1080
gcgcctccac ctccagatata tgacaagcag ttggatgaaa gagaacacac aattgaagaa	1140
tggaaagaac ttatctacaa ggaagtaatg aattcagaag aaaagactaa aaatggtgta	1200
gtaaaaggac agccttctcc ttcagcacag gtgcagcagt gaacagcagt gagagtctcc	1260
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cctgcctgcg aaaccacgcg ttcttcagga gatgatgtga tggaacacac acacacgcag	1440
acacacacac acacacaaat gcagacacac aacatcaaga aaacagcaag ggagagaatc	1500
caagcctaaa attaaataaa tctttcagcc tgcttcttcc ccagggttct gtattgcagc	1560
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ttactacaga aagcaaatca gacacaatta gagaagcctt ttccataaag tgtaatttta	1680
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ggcccatcc agcatgtgtg tgtctctatc ttgcatctac ctgctccttg gcctagtcag	1800
atggatgtag atacagatcc gcatgtgtct gtattcatac agcactactt acttagagat	1860
gctactgtca gtgtcctcag ggctctacca agacataatg cactggggta ccacatggtc	1920
catttcatgt gatctattac tctgacataa acccatctgt aatatattgc cagtatataa	1980
gctgtttagt ttgttaattg attaaactgt atgtcttata agaaaacatg taaaggggga	2040
atatatgggg ggagtgcgct ctctcagacc cttgaagatg tagcttccaa atttgaatgg	2100

attaaatggc acctgtatac caatttgtag a

2131

<210> 88
<211> 989
<212> DNA
<213> Homo sapiens

<400> 88
gagaaatggc gtggcagggg acccagcgag cccagagggg ttttgccgct gcttcctcta 60
cccctgtatt tcacgcagct ctctaaattg actcagctcc aggctagtgt gagaaacacc 120
aacagcagggc ccatctcaga tcttcactat ggcaacttat gcaagaaaact gttgaattag 180
accggtttcc tatagatgag aaaccataca agctgtggta tttatgagcc tccatttctt 240
atactactgc agtgaaccaa cattggatgt gaaaattgcc tttgttcagg gattcgataa 300
acaagtggat gtgtcatata ttgccaaaca ttacaacatg agcaaaaagca aagttgacaa 360
ccagttctac agtgtggaag tgggagactc aaccttcaca gttctcaagc gctaccagaa 420
tctaaagcct attggctctg gggctcaggg catagtttgt gccgcgtatg atgctgtcct 480
tgacagaaat gtggccatta agaagctcag cagacccttt cagaaccaaa cacatgccaa 540
gagagcgtac cgggagctgg tcctcatgaa gtgtgtgaac cataaaaacg tgagttttgt 600
tatttttaaa cttctggcag tgggagtttg taagattgga aaaagaaaat gtgtctgtac 660
ttgagtaaga aaggctttct ttgctttgca attcctgcct cttaagattg tttatcctgc 720
caatatgcta cattcgattt cattgtcctc atggtagctt tctgcttaaa aatcacctaa 780
aaaccatatg atgtggaagt gatgaatttt ataaatattt ccctagttag aaaatttacc 840
atatttggtt atcttgctcc aaattgagag cttcaagaaa gaaacaagac aaacaaagcc 900
caaagccaga aaaaaaaaaa aacctacaaa gcaccatgct acatttcttt ataatttaga 960
tttaatttag aatttattaa tttaaaatt 989

<210> 89
<211> 1818
<212> DNA
<213> Homo sapiens

<400> 89
gaattccggg ccaggcatgg tagcgcacgc ctgtaatccc agctactcgg gaaactgagg 60
tgggagaatc gattgaacct ggaagtggag gttgcggtga gccaaagatca tcctgtcgca 120
ctccagcctg ggcaacaaga gcgaaactcc atctcaaaaa gaaaaaaaaa gatatatatg 180
tgtgacttac aggtacaggt aaagttgctt ctggttttct gggtgttgca tggattttcc 240

tatgcagcca caggtcttta ttttcttact taagtgcctc caacttccca taacacaaat	300
taaggcatga tgaacatcct ctctgtgctg aacatcctgt gtatgtcact tcagaagcct	360
gtgtgacggg ttcttttagtc tttataccta ggggtgggat ttctgggtca taggacagta	420
atttatattt atttcactaa gtattctctt tctctggctt ttgttacata ttacctgttt	480
gtcctccaga aaacttgcac caatttacat tcctaccaat agggtaggag agtgcacaat	540
gggtggattc taactccaaa tctaacacct cttcttttct ttgtttctag cagccatggc	600
aatgacaggc tcaacacctt gctcatccat gagtaaccac acaaaggaaa gggtgacaat	660
gaccaaagtg aactggaga atttttatag caaccttacc gctcaacatg aagaacgaga	720
aatgagacaa aagaagttag aaaagggtgat ggaagaagaa ggcctaaaag atgaggagaa	780
acgactccgg agatcagcac atgctcggaa ggaaacagag tttcttcgtt tgaagagaac	840
aagacttgga ttggaagatt ttgagtcctt aaaagtaata ggcagaggag catttggtga	900
ggtaaggctt gttcagaaga aagatacggg acatgtgtat gcaatgaaaa tactccgtaa	960
agcagatatg cttgaaaaag agcagggttg ccacattcgt gcggagcgtg acattctagt	1020
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cactctgacg aagaggagac tcagttttat atagcagaaa cagtattagc catagactct	1200
attcaccaac ttggattcat ccacagagac atcaaaccag acaaccttct tttggacagc	1260
aagggccatg tgaaactttc tgactttggt ctttgcacag gactgaaaaa agcacatagg	1320
acagaatttt ataggaatct gaaccacagc ctccccagtg atttcacttt ccagaacatg	1380
aattccaaaa ggaaagcaga aacctggaaa agaaatagac gtcagctagc cttctccaca	1440
gtaggcactc ctgactacat tgctcctgag gtgttcatgc agaccgggta caacaagctc	1500
tgtgattggg ggtcgcttg ggtgatcatg tatgagatgc tcatcggtaa gttgcatggg	1560
ttcagaggac tttttctgtg catccatgac agacttttac attgatacca gcctctgttt	1620
caattggcag tgatctaagt gatctcccta cttgtcttcc aaagtgaatt gttttagaca	1680
gatgacacct ctttcagtaa gatgtatccc actccattct tgggcttact ggcatcctgc	1740
aattgctttg ctgatcattt tttatgtttt tctttctctt ataccttcat cttctccatc	1800
tagaagctct tttagtca	1818

<211> 2732
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(2732)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 90
ggcacgancg gcacgagtcg gcacgagcgg cacgagtaag tggagggttca atgttggata 60
tcataaaata cattgtcaac cgaggagaac acaagaatgg agttctggaa gaggcaataa 120
tagcaacaat tottaaagag gttttggaag gcttagacta tctacacaga aacggtcaga 180
ttcacaggga tttgaaagct ggtaatatc ttctgggtga ggatgggttca gtacaaatag 240
cagattttgg ggtaagtgcg ttcttagcaa caggggggtga tgttaccoga aataaagtaa 300
gaaaaacatt cgttggcacc ccatgttggga tggctcctga agtcatggaa caggtgagag 360
gctatgactt caaggctgac atgtggagtt ttggaataac tgccattgaa ttagcaacag 420
gagcagcgcc ttatcacaaa tctctccca tgaaagtgtt aatgttgact ttgcaaaatg 480
atccaccac tttggaaaca ggggtagagg ataaagaaat gatgaaaaag tacggcaagt 540
cctttagaaa attactttca ctgtgtcttc agaaagatcc ttccaaaagg cccacagcag 600
cagaactttt aaaatgcaaa ttcttcaga aagccaagaa cagagagtac ctgattgaga 660
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ggccaagtgg tcaccttcat aaaaccgaag acggggactg ggagtggagt gacgacgaga 780
tggatgagaa gagcgaagaa gggaaagcag ctttttctca ggaaaagtca cgaagagtaa 840
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gatttgagtt tactccagga agagatacag cagatgggtg atctcaggag ctcttctctg 1080
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agattcctga tgaagtgaag ctgattgggt ttgctcagtt gagtgtcagc tgatgtatgt 1260
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tagcaacaat tcttaaagag gttttggaag gcttagacta tctacacaga aacggtcaga 180
ttcacaggga tttgaaagct ggtaatatc ttctgggtga ggatgggttca gtacaaatag 240
cagatTTTgg ggtaagtgcg ttcctagcaa caggggggtga tgttaccga aataaagtaa 300
gaaaaacatt cgttggcacc ccatgttgga tggctcctga agtcatggaa caggtgagag 360
gctatgactt caaggctgac atgtggagtt ttggaataac tgccattgaa ttagcaacag 420
gagcagcgc ttatcacaaa tctctccca tgaaagtgtt aatgttgact ttgcaaaatg 480
atcccccac tttgaaaca ggggtagagg ataaagaaat gatgaaaaag tacggcaagt 540
cctttagaaa attactttca ctgtgtcttc agaaagatcc ttccaaaagg cccacagcag 600
cagaactttt aaatgcaaa ttcttcaga aagccaagaa cagagagtac ctgattgaga 660
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ggtcaagtgg tcaccttcac aaaaccgaag acggggactg ggagtggagt gacgacgaga 780
tggatgagaa gagcgaagaa gggaaagcag ctttttctca ggaaaagtca cgaagagtaa 840
aagaagaaaa tccagagatt gcagtgagtg ccagcaccat cccgaacaa atacagtccc 900
tctctgtgca cgactctcag ggccccacca atgctaataga agactacaga gaagcttctt 960
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ctgaattcaa agcaaacactc ttaataaagc tttcct 1416

<210> 92
<211> 434
<212> PRT
<213> Homosapiens

<400> 92

Met 1	Pro	Ala	Arg	Arg 5	Leu	Leu	Leu	Leu	Leu 10	Thr	Leu	Leu	Leu	Pro 15	Gly
Leu	Gly	Ile	Phe 20	Gly	Ser	Thr	Ser	Thr 25	Val	Thr	Leu	Pro	Glu 30	Thr	Leu
Leu	Phe 35	Val	Ser	Thr	Leu	Asp	Gly 40	Ser	Leu	His	Ala	Val 45	Ser	Lys	Arg
Thr	Gly 50	Ser	Ile	Lys	Trp	Thr 55	Leu	Lys	Glu	Asp 60	Pro	Val	Leu	Gln	Val
Pro 65	Thr	His	Val	Glu 70	Glu	Pro	Ala	Phe	Leu 75	Pro	Asp	Pro	Asn	Asp 80	Gly
Ser	Leu	Tyr	Thr 85	Leu	Gly	Ser	Lys	Asn 90	Asn	Glu	Gly	Leu	Thr 95	Lys	Leu
Pro	Phe	Thr	Ile 100	Pro	Glu	Leu	Val	Gln 105	Ala	Ser	Pro	Cys	Arg 110	Ser	Ser
Asp	Gly 115	Ile	Leu	Tyr	Met	Gly	Lys 120	Lys	Gln	Asp	Ile	Trp 125	Tyr	Val	Ile
Asp 130	Leu	Leu	Thr	Gly	Glu	Lys 135	Gln	Gln	Thr	Leu 140	Ser	Ser	Ala	Phe	Ala
Asp 145	Ser	Leu	Cys	Pro 150	Ser	Thr	Ser	Leu	Leu	Tyr 155	Leu	Gly	Arg	Thr	Glu
Tyr	Thr	Ile	Thr 165	Met	Tyr	Asp	Thr	Lys 170	Thr	Arg	Glu	Leu	Arg 175	Trp	Asn
Ala	Thr	Tyr	Phe 180	Asp	Tyr	Ala	Ala	Ser 185	Leu	Pro	Glu	Asp 190	Glu	Gly	Asp
Tyr	Lys 195	Met	Ser	His	Phe	Val	Ser 200	Asn	Gly	Asp	Gly	Leu 205	Val	Val	Thr
Val 210	Asp	Ser	Glu	Ser	Gly	Asp 215	Val	Leu	Trp	Ile	Gln 220	Asn	Tyr	Ala	Ser
Pro 225	Val	Val	Ala	Phe 230	Tyr	Val	Trp	Gln	Arg	Glu 235	Gly	Leu	Arg	Lys	Val
Met	His	Ile	Asn 245	Val	Ala	Val	Glu	Thr	Leu 250	Arg	Tyr	Leu	Thr 255	Phe	Met
Ser	Gly	Glu	Val 260	Gly	Arg	Ile	Thr	Lys 265	Trp	Lys	Tyr	Pro	Phe 270	Pro	Lys
Glu	Thr 275	Glu	Ala	Lys	Ser	Lys	Leu 280	Thr	Pro	Thr	Leu	Tyr 285	Val	Gly	Lys
Tyr 290	Ser	Thr	Ser	Leu	Tyr	Ala 295	Ser	Pro	Ser	Met	Val 300	His	Glu	Gly	Val

Ala Val Val Pro Arg Gly Ser Thr Leu Pro Leu Leu Glu Gly Pro Gln
305 310 315 320

Thr Asp Gly Val Thr Ile Gly Asp Lys Gly Glu Cys Val Ile Thr Pro
325 330 335

Ser Thr Asp Val Lys Phe Asp Pro Gly Leu Lys Ser Lys Asn Lys Leu
340 345 350

Asn Tyr Leu Arg Asn Tyr Trp Leu Leu Ile Gly His His Glu Thr Pro
355 360 365

Leu Ser Ala Ser Thr Lys Met Leu Glu Arg Phe Pro Asn Asn Leu Pro
370 375 380

Lys His Arg Glu Asn Val Ile Pro Ala Asp Ser Glu Lys Lys Ser Phe
385 390 395 400

Glu Glu Thr Leu Leu Gln Met Thr Ser Val Phe Ser Trp Ile Leu Asn
405 410 415

Leu Pro Ser Lys Glu Glu Val Phe Ala Phe Leu Arg Ile Phe Glu Lys
420 425 430

Asn Met

<210> 93
<211> 232
<212> PRT
<213> Homo sapiens

<400> 93

Met Tyr Ser Leu Gln Leu Gln Ser Val Ser Ser Ala Ile His Leu Cys
1 5 10 15

Asp Lys Lys Lys Met Glu Leu Ser Leu Asn Ile Pro Val Asn His Gly
20 25 30

Pro Gln Glu Glu Ser Cys Gly Ser Ser Gln Leu His Glu Asn Ser Gly
35 40 45

Ser Pro Glu Thr Ser Arg Ser Leu Pro Ala Pro Gln Asp Asn Asp Phe
50 55 60

Leu Ser Arg Lys Ala Gln Asp Cys Tyr Phe Met Lys Leu His His Cys
65 70 75 80

Pro Gly Asn His Ser Trp Asp Ser Thr Ile Ser Gly Ser Gln Arg Ala
85 90 95

Ala Phe Cys Asp His Lys Thr Thr Pro Cys Ser Ser Ala Ile Ile Asn
100 105 110

Pro Leu Ser Thr Ala Gly Asn Ser Glu Arg Leu Gln Pro Gly Ile Ala

007460404

115					120					125					
Gln	Gln	Trp	Ile	Gln	Ser	Lys	Arg	Glu	Asp	Ile	Val	Asn	Gln	Met	Thr
130						135					140				
Glu	Ala	Cys	Leu	Asn	Gln	Ser	Leu	Asp	Ala	Leu	Leu	Ser	Arg	Asp	Leu
145						150					155				160
Ile	Met	Lys	Glu	Asp	Tyr	Glu	Leu	Val	Ser	Thr	Lys	Pro	Thr	Arg	Thr
				165					170					175	
Ser	Lys	Val	Arg	Gln	Leu	Leu	Asp	Thr	Thr	Asp	Ile	Gln	Gly	Glu	Glu
			180					185					190		
Phe	Ala	Lys	Val	Ile	Val	Gln	Lys	Leu	Lys	Asp	Asn	Lys	Gln	Met	Gly
		195					200					205			
Leu	Gln	Pro	Tyr	Pro	Glu	Ile	Leu	Val	Val	Ser	Arg	Ser	Pro	Ser	Leu
	210					215					220				
Asn	Leu	Leu	Gln	Asn	Lys	Ser	Met								
225						230									
<210>	94														
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<212>	PRT														
<213>	Homo sapiens														
<400>	94														
Met	Ala	Asp	Leu	Glu	Ala	Val	Leu	Ala	Asp	Val	Ser	Tyr	Leu	Met	Ala
1				5					10					15	
Met	Glu	Lys	Ser	Lys	Ala	Thr	Pro	Ala	Ala	Arg	Ala	Ser	Lys	Lys	Ile
			20					25					30		
Leu	Leu	Pro	Glu	Pro	Ser	Ile	Arg	Ser	Val	Met	Gln	Lys	Tyr	Leu	Glu
		35					40					45			
Asp	Arg	Gly	Glu	Val	Thr	Phe	Glu	Lys	Ile	Phe	Ser	Gln	Lys	Leu	Gly
	50					55					60				
Tyr	Leu	Leu	Phe	Arg	Asp	Phe	Cys	Leu	Asn	His	Leu	Glu	Glu	Ala	Arg
65				70							75				80
Pro	Leu	Val	Glu	Phe	Tyr	Glu	Glu	Ile	Lys	Lys	Tyr	Glu	Lys	Leu	Glu
				85				90						95	
Thr	Glu	Glu	Glu	Arg	Val	Ala	Arg	Ser	Arg	Glu	Ile	Phe	Asp	Ser	Tyr
			100					105					110		
Ile	Met	Lys	Glu	Leu	Leu	Ala	Cys	Ser	His	Pro	Phe	Ser	Lys	Ser	Ala
		115					120					125			
Thr	Glu	His	Val	Gln	Gly	His	Leu	Gly	Lys	Lys	Gln	Val	Pro	Pro	Asp
	130					135					140				

115 120 125
 130 135 140
 145 150 155 160
 165 170 175
 180 185 190
 195 200 205
 210 215 220
 225 230
 <210> 94
 <211> 209
 <212> PRT
 <213> Homo sapiens
 <400> 94
 1 5 10 15
 20 25 30
 35 40 45
 50 55 60
 65 70 75 80
 85 90 95
 100 105 110
 115 120 125
 130 135 140

Leu Phe Gln Pro Tyr Ile Glu Glu Ile Cys Gln Asn Leu Arg Gly Asp
145 150 155 160

Val Phe Gln Lys Phe Ile Glu Arg Val Ala Leu Ala Ala Gly Ala Ala
165 170 175

Thr Leu Pro Ala Val Pro Ser Cys Pro Asn Pro Gln His Pro Gly Ser
180 185 190

Gly Thr Thr Ala Arg His Leu Gln Val Gly Pro Tyr Trp Pro Arg Leu
195 200 205

Ala

<210> 95
<211> 454
<212> PRT
<213> Homo sapiens

<400> 95

Met Gly Leu Val Ser Ser Lys Lys Pro Asp Lys Glu Lys Pro Ile Lys
1 5 10 15

Glu Lys Asp Lys Gly Gln Trp Ser Pro Leu Lys Val Ser Ala Gln Asp
20 25 30

Lys Asp Ala Pro Pro Leu Pro Pro Leu Val Val Phe Asn His Leu Thr
35 40 45

Pro Pro Pro Pro Asp Glu His Leu Asp Glu Asp Lys His Phe Val Val
50 55 60

Ala Leu Tyr Asp Tyr Thr Ala Met Asn Asp Arg Asp Leu Gln Met Leu
65 70 75 80

Lys Gly Glu Lys Leu Gln Val Leu Lys Gly Thr Gly Asp Trp Trp Leu
85 90 95

Ala Arg Ser Leu Val Thr Gly Arg Glu Gly Tyr Val Pro Ser Asn Phe
100 105 110

Val Ala Arg Val Glu Ser Leu Glu Met Glu Arg Trp Phe Phe Arg Ser
115 120 125

Gln Gly Arg Lys Glu Ala Glu Arg Gln Leu Leu Ala Pro Ile Asn Lys
130 135 140

Ala Gly Ser Phe Leu Ile Arg Glu Ser Glu Thr Asn Lys Gly Ala Phe
145 150 155 160

Ser Leu Ser Val Lys Asp Val Thr Thr Gln Gly Glu Leu Ile Lys His
165 170 175

Tyr Lys Ile Arg Cys Leu Asp Glu Gly Gly Tyr Tyr Ile Ser Pro Arg
180 185 190

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Ile Thr Phe Pro Ser Leu Gln Ala Leu Val Gln His Tyr Ser Ser Tyr
 195 200 205

Tyr Lys Asn Asn Met Lys Val Ala Ile Lys Thr Leu Lys Glu Gly Thr
 210 215 220

Met Ser Pro Glu Ala Phe Leu Gly Glu Ala Asn Val Met Lys Ala Leu
 225 230 235 240

Gln His Glu Arg Leu Val Arg Leu Tyr Ala Val Val Thr Lys Glu Pro
 245 250 255

Ile Tyr Ile Val Thr Glu Tyr Met Ala Arg Gly Cys Leu Leu Asp Phe
 260 265 270

Leu Lys Thr Asp Glu Gly Ser Arg Leu Ser Leu Pro Arg Leu Ile Asp
 275 280 285

Met Ser Ala Gln Ile Ala Glu Gly Met Ala Tyr Ile Glu Arg Met Asn
 290 295 300

Ser Ile His Arg Asp Leu Arg Ala Ala Asn Ile Leu Val Ser Glu Ala
 305 310 315 320

Leu Cys Cys Lys Ile Ala Asp Phe Gly Leu Ala Arg Ile Ile Asp Ser
 325 330 335

Glu Tyr Thr Ala Gln Glu Gly Ala Lys Phe Pro Ile Lys Trp Thr Ala
 340 345 350

Pro Glu Ala Ile His Phe Gly Val Phe Thr Ile Lys Ala Asp Val Trp
 355 360 365

Ser Phe Gly Val Leu Leu Met Glu Val Val Thr Tyr Gly Arg Val Pro
 370 375 380

Tyr Pro Gly Met Ser Asn Pro Glu Val Ile Arg Asn Leu Glu Arg Gly
 385 390 395 400

Tyr Arg Met Pro Arg Pro Asp Thr Cys Pro Pro Glu Leu Tyr Arg Gly
 405 410 415

Val Ile Ala Glu Cys Trp Arg Ser Arg Pro Glu Glu Arg Pro Thr Phe
 420 425 430

Glu Phe Leu Gln Ser Val Leu Glu Asp Phe Tyr Thr Ala Thr Glu Arg
 435 440 445

Gln Tyr Glu Leu Gln Pro
 450

<210> 96
 <211> 82
 <212> PRT
 <213> Homo sapiens

<400> 96

Met Glu Asn Phe Gln Lys Val Glu Lys Ile Gly Glu Gly Thr Tyr Gly
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Val Val Tyr Lys Ala Arg Asn Lys Leu Thr Gly Glu Val Val Ala Leu
20 25 30
Lys Lys Ile Arg Leu Asp Thr Glu Thr Glu Gly Val Pro Ser Thr Ala
35 40 45
Ile Arg Glu Ile Ser Leu Leu Lys Glu Leu Asn His Pro Asn Ile Val
50 55 60
Lys Leu Leu Asp Val Ile His Thr Glu Asn Lys Asn Ile Ser Leu Lys
65 70 75 80
Glu Gly

<210> 97

<211> 118

<212> PRT

<213> Homo sapiens

<400> 97

Met Thr Arg Asp Glu Ala Leu Pro Asp Ser His Ser Ala Gln Asp Phe
1 5 10 15
Tyr Glu Asn Tyr Glu Pro Lys Glu Ile Leu Gly Arg Gly Val Ser Ser
20 25 30
Val Val Arg Arg Cys Ile His Lys Pro Thr Ser Gln Glu Tyr Ala Val
35 40 45
Lys Val Ile Asp Val Thr Gly Gly Gly Ser Phe Ser Pro Glu Glu Val
50 55 60
Arg Glu Leu Arg Glu Ala Thr Leu Lys Glu Val Asp Ile Leu Arg Lys
65 70 75 80
Val Ser Gly His Pro Asn Ile Ser Ile Gln Leu Lys Asp Thr Tyr Glu
85 90 95
Thr Asn Thr Phe Phe Phe Leu Val Phe Asp Leu Met Lys Arg Gly Glu
100 105 110
Leu Phe Asp Leu Pro His
115

<210> 98

<211> 167

<212> PRT

<213> Homo sapiens

<400> 98

Val Phe Leu Gly Arg Cys Arg Ser Val Lys Glu Phe Glu Lys Leu Asn
 1 5 10 15
 Arg Ile Gly Glu Gly Thr Tyr Gly Ile Val Tyr Arg Ala Arg Asp Thr
 20 25 30
 Gln Thr Asp Glu Ile Val Ala Leu Lys Lys Val Arg Met Asp Lys Glu
 35 40 45
 Lys Asp Gly Ile Pro Ile Ser Ser Leu Arg Glu Ile Thr Leu Leu Leu
 50 55 60
 Arg Leu Arg His Pro Asn Ile Val Glu Leu Lys Glu Val Val Val Gly
 65 70 75 80
 Asn His Leu Glu Ser Ile Phe Leu Val Met Gly Tyr Cys Glu Gln Asp
 85 90 95
 Leu Ala Ser Leu Leu Glu Asn Met Pro Thr Pro Phe Ser Glu Ala Gln
 100 105 110
 Val Lys Cys Ile Val Leu Gln Val Leu Arg Gly Leu Gln Tyr Leu His
 115 120 125
 Arg Asn Phe Ile Ile His Arg Asp Leu Lys Val Ser Asn Leu Leu Met
 130 135 140
 Thr Asp Lys Gly Cys Val Lys Thr Gly Gly Cys Asn Leu Gly Gln Ala
 145 150 155 160
 Trp Ser Leu Asp Gly Thr Trp
 165

<210> 99
 <211> 141
 <212> PRT
 <213> Homo sapiens

<400> 99

Met Ser Ser Ala Gly Gly Val Ser Arg Arg Leu Ala Ala Val Arg Ser
 1 5 10 15
 Thr Val Leu Cys Arg Ala Val Gly Cys Ile Leu Ala Glu Leu Leu Ala
 20 25 30
 His Arg Pro Leu Leu Pro Gly Thr Ser Glu Ile His Gln Ile Asp Leu
 35 40 45
 Ile Val Gln Leu Leu Gly Thr Pro Ser Glu Asn Ile Trp Pro Gly Phe
 50 55 60
 Ser Lys Leu Pro Leu Val Gly Gln Tyr Ser Leu Arg Lys Gln Pro Tyr
 65 70 75 80
 Asn Asn Leu Lys His Lys Phe Pro Trp Leu Ser Glu Ala Gly Leu Arg

			85					90					95						
Leu	Leu	His	Phe	Leu	Phe	Met	Tyr	Asp	Pro	Lys	Lys	Arg	Ala	Thr	Ala				
			100					105					110						
Gly	Asp	Cys	Leu	Glu	Ser	Ser	Tyr	Phe	Lys	Glu	Lys	Pro	Leu	Arg	Leu				
		115					120					125							
Pro	Ile	Ser	Gly	Val	Cys	Glu	Gly	Cys	Arg	Glu	Pro	Gly							
	130					135					140								

<210> 100
 <211> 119
 <212> PRT
 <213> Homo sapiens

<400> 100

Val	Phe	Leu	Gly	Arg	Cys	Arg	Ser	Val	Lys	Glu	Phe	Glu	Lys	Leu	Asn				
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Arg	Ile	Gly	Glu	Gly	Thr	Tyr	Gly	Ile	Val	Tyr	Arg	Ala	Arg	Asp	Thr				
		20					25						30						
Gln	Thr	Asp	Glu	Ile	Val	Ala	Leu	Lys	Lys	Val	Arg	Met	Asp	Lys	Glu				
		35					40					45							
Lys	Asp	Gly	Ile	Pro	Ile	Ser	Ser	Leu	Arg	Glu	Ile	Thr	Leu	Leu	Leu				
	50					55					60								
Arg	Leu	Arg	His	Pro	Asn	Ile	Leu	Pro	Ala	Arg	Ala	Pro	Trp	Lys	Gly				
	65				70					75					80				
Arg	Ser	Gly	Gly	Ser	Ile	Arg	Gly	Cys	Arg	Gly	Leu	Met	Trp	Ser	Ser				
				85				90						95					
Ser	Leu	Cys	Trp	Lys	Cys	Ala	Thr	Thr	Ala	Ser	Trp	Glu	Glu	Trp	Trp				
		100					105						110						

Val	Gln	Ser	Pro	Arg	Cys	Leu													
		115																	

<210> 101
 <211> 756
 <212> PRT
 <213> Homo sapiens

<400> 101

Met	Gly	Glu	Ala	Glu	Lys	Phe	His	Tyr	Ile	Tyr	Ser	Cys	Asp	Leu	Asp				
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Ile	Asn	Val	Gln	Leu	Lys	Ile	Gly	Ser	Leu	Glu	Gly	Lys	Arg	Glu	Gln				
		20					25						30						
Lys	Ser	Tyr	Lys	Ala	Val	Leu	Glu	Asp	Pro	Met	Leu	Lys	Phe	Ser	Gly				
		35					40					45							

Leu Tyr Gln Glu Thr Cys Ser Asp Leu Tyr Val Thr Cys Gln Val Phe
 50 55 60
 Ala Glu Gly Lys Pro Leu Ala Leu Pro Val Arg Thr Ser Tyr Lys Ala
 65 70 75 80
 Phe Ser Thr Arg Trp Asn Trp Asn Glu Trp Leu Lys Leu Pro Val Lys
 85 90 95
 Tyr Pro Asp Leu Pro Arg Asn Ala Gln Val Ala Leu Thr Ile Trp Asp
 100 105 110
 Val Tyr Gly Pro Gly Lys Ala Val Pro Val Gly Gly Thr Thr Val Ser
 115 120 125
 Leu Phe Gly Lys Tyr Gly Met Phe Arg Gln Gly Met His Asp Leu Lys
 130 135 140
 Val Trp Pro Asn Val Glu Ala Asp Gly Ser Glu Pro Thr Lys Thr Pro
 145 150 155 160
 Gly Arg Thr Ser Ser Thr Leu Ser Glu Asp Gln Met Ser Arg Leu Ala
 165 170 175
 Lys Leu Thr Lys Ala His Arg Gln Gly His Met Val Lys Val Asp Trp
 180 185 190
 Leu Asp Arg Leu Thr Phe Arg Glu Ile Glu Met Ile Asn Glu Ser Val
 195 200 205
 Lys Arg Ser Ser Asn Phe Met Tyr Leu Met Gly Gly Phe Arg Cys Val
 210 215 220
 Lys Cys Asp Asp Lys Glu Tyr Gly Ile Val Tyr Tyr Glu Lys Asp Gly
 225 230 235 240
 Asp Glu Ser Ser Pro Ile Leu Thr Ser Phe Glu Leu Val Lys Val Pro
 245 250 255
 Asp Pro Gln Met Ser Leu Glu Asn Leu Val Glu Ser Lys His His Asn
 260 265 270
 Leu Pro Arg Ser Leu Arg Ser Gly Pro Ser Asp His Asp Leu Lys Pro
 275 280 285
 Tyr Pro Ser Pro Arg Asp Gln Leu Lys Asn Ile Val Ser Tyr Pro Pro
 290 295 300
 Ser Lys Pro Pro Thr Tyr Glu Glu Gln Asp Leu Val Trp Glu Phe Arg
 305 310 315 320
 Tyr Tyr Leu Thr Asn Gln Asp Lys Ala Leu Thr Lys Ile Leu Thr Ser
 325 330 335
 Val Ile Trp Asp Leu Pro Gln Glu Ala Lys Gln Ala Leu Ala Leu Leu
 340 345 350

Gly Lys Trp Asn Pro Met Asp Val Glu Asp Ser Leu Glu Leu Ile Ser
355 360 365

Ser His Tyr Thr Asn Pro Thr Val Arg Arg Tyr Ala Val Ala Arg Leu
370 375 380

Arg Gln Ala Asp Asp Glu Asp Leu Leu Met Tyr Leu Leu Gln Leu Val
385 390 395 400

Gln Ala Leu Lys Tyr Glu Asn Phe Asp Asp Ile Lys Asn Gly Leu Glu
405 410 415

Pro Thr Lys Lys Asp Ser Gln Ser Ser Val Ser Glu Asn Val Ser Asn
420 425 430

Ser Gly Ile Asn Ser Ala Glu Ile Asp Ser Ser Gln Ile Ile Thr Ser
435 440 445

Pro Leu Pro Ser Val Ser Ser Pro Pro Pro Ala Ser Lys Thr Lys Glu
450 455 460

Val Pro Asp Gly Glu Asn Leu Glu Gln Asp Leu Cys Thr Phe Leu Ile
465 470 475 480

Ser Arg Ala Cys Lys Asn Ser Thr Leu Ala Asn Tyr Leu Tyr Trp Tyr
485 490 495

Val Ile Val Glu Cys Glu Asp Gln Asp Thr Gln Gln Arg Asp Pro Lys
500 505 510

Thr His Glu Met Tyr Leu Asn Val Met Arg Arg Phe Ser Gln Ala Leu
515 520 525

Leu Lys Gly Asp Lys Ser Val Arg Val Met Arg Ser Leu Leu Ala Ala
530 535 540

Gln Gln Thr Phe Val Asp Arg Leu Val His Leu Met Lys Ala Val Gln
545 550 555 560

Arg Glu Ser Gly Asn Arg Lys Lys Lys Asn Glu Arg Leu Gln Ala Leu
565 570 575

Leu Gly Asp Asn Glu Lys Met Asn Leu Ser Asp Val Glu Leu Ile Pro
580 585 590

Leu Pro Leu Glu Pro Gln Val Lys Ile Arg Gly Ile Ile Pro Glu Thr
595 600 605

Ala Thr Leu Phe Lys Ser Ala Leu Met Pro Ala Gln Leu Phe Phe Lys
610 615 620

Thr Glu Asp Gly Gly Lys Tyr Pro Val Ile Phe Lys His Gly Asp Asp
625 630 635 640

Leu Arg Gln Asp Gln Leu Ile Leu Gln Ile Ile Ser Leu Met Asp Lys
645 650 655

Leu Leu Arg Lys Glu Asn Leu Asp Leu Lys Leu Thr Pro Tyr Lys Val
660 665 670

Leu Ala Thr Ser Thr Lys His Gly Phe Met Gln Phe Ile Gln Ser Val
675 680 685

Pro Val Ala Glu Val Leu Asp Thr Glu Gly Ser Ile Gln Asn Phe Phe
690 695 700

Arg Lys Tyr Ala Pro Ser Glu Asn Gly Pro Asn Gly Ile Ser Ala Glu
705 710 715 720

Val Met Asp Thr Tyr Val Lys Ser Cys Ala Gly Tyr Cys Val Ile Thr
725 730 735

Tyr Ile Leu Gly Val Gly Asp Arg His Leu Asp Asn Leu Leu Leu Thr
740 745 750

Lys Thr Gly Gly
755

<210> 102
<211> 508
<212> PRT
<213> Homo sapiens

<400> 102

Met Gly Glu Ala Glu Lys Phe His Tyr Ile Tyr Ser Cys Asp Leu Asp
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Ile Asn Val Gln Leu Lys Ile Gly Ser Leu Glu Gly Lys Arg Glu Gln
20 25 30

Lys Ser Tyr Lys Ala Val Leu Glu Asp Pro Met Leu Lys Phe Ser Gly
35 40 45

Leu Tyr Gln Glu Thr Cys Ser Asp Leu Tyr Val Thr Cys Gln Val Phe
50 55 60

Ala Glu Gly Lys Pro Leu Ala Leu Pro Val Arg Thr Ser Tyr Lys Ala
65 70 75 80

Phe Ser Thr Arg Trp Asn Trp Asn Glu Trp Leu Lys Leu Pro Val Lys
85 90 95

Tyr Pro Asp Leu Pro Arg Asn Ala Gln Val Ala Leu Thr Ile Trp Asp
100 105 110

Val Tyr Gly Pro Gly Lys Ala Val Pro Val Gly Gly Thr Thr Val Ser
115 120 125

Leu Phe Gly Lys Tyr Gly Met Phe Arg Gln Gly Met His Asp Leu Lys
130 135 140

Val Trp Pro Asn Val Glu Ala Asp Gly Ser Glu Pro Thr Lys Thr Pro

145		150		155		160
Gly Arg Thr Ser Ser Thr Leu Ser Glu Asp Gln Met Ser Arg Leu Ala						
	165			170		175
Lys Leu Thr Lys Ala His Arg Gln Gly His Met Val Lys Val Asp Trp						
	180			185		190
Leu Asp Arg Leu Thr Phe Arg Glu Ile Glu Met Ile Asn Glu Ser Val						
	195			200		205
Lys Arg Ser Ser Asn Phe Met Tyr Leu Met Gly Gly Phe Arg Cys Val						
	210			215		220
Lys Cys Asp Asp Lys Glu Tyr Gly Ile Val Tyr Tyr Glu Lys Asp Gly						
	225			230		235
Asp Glu Ser Ser Pro Ile Leu Thr Ser Phe Glu Leu Val Lys Val Pro						
	245			250		255
Asp Pro Gln Met Ser Leu Glu Asn Leu Val Glu Ser Lys His His Asn						
	260			265		270
Leu Pro Arg Ser Leu Arg Ser Gly Pro Ser Asp His Asp Leu Lys Pro						
	275			280		285
Tyr Pro Ser Pro Arg Asp Gln Leu Lys Asn Ile Val Ser Tyr Pro Pro						
	290			295		300
Ser Lys Pro Pro Thr Tyr Glu Glu Gln Asp Leu Val Trp Glu Phe Arg						
	305			310		315
Tyr Tyr Leu Thr Asn Gln Asp Lys Ala Leu Thr Lys Ile Leu Thr Ser						
	325			330		335
Val Ile Trp Asp Leu Pro Gln Glu Ala Lys Gln Ala Leu Ala Leu Leu						
	340			345		350
Gly Lys Trp Asn Pro Met Asp Val Glu Asp Ser Leu Glu Leu Ile Ser						
	355			360		365
Ser His Tyr Thr Asn Pro Thr Val Arg Arg Tyr Ala Val Ala Arg Leu						
	370			375		380
Arg Gln Ala Asp Asp Glu Asp Leu Leu Met Tyr Leu Leu Gln Leu Val						
	385			390		395
Gln Ala Leu Lys Tyr Glu Asn Phe Asp Asp Ile Lys Asn Gly Leu Glu						
	405			410		415
Pro Thr Lys Lys Asp Ser Gln Ser Ser Val Ser Glu Asn Val Ser Asn						
	420			425		430
Ser Gly Ile Asn Ser Ala Glu Ile Asp Ser Ser Gln Ile Ile Thr Ser						
	435			440		445
Pro Leu Pro Ser Val Ser Ser Pro Pro Pro Ala Ser Lys Thr Lys Glu						

145 150 155 160

450 455 460
 Val Pro Asp Gly Glu Asn Leu Glu Gln Asp Leu Cys Thr Phe Leu Ile
 465 470 475 480
 Ser Arg Ala Cys Lys Asn Ser Thr Leu Ala Asn Tyr Leu Tyr Trp Tyr
 485 490 495
 Val Lys Ile Ile Phe Cys Leu Phe Ser Tyr Tyr Pro
 500 505

<210> 103
 <211> 140
 <212> PRT
 <213> Homo sapiens

<400> 103

Met Gly Asn Ala Ala Ala Ala Lys Lys Gly Ser Glu Gln Glu Ser Val
 1 5 10 15
 Lys Glu Phe Leu Ala Lys Ala Lys Glu Asp Phe Leu Lys Lys Trp Glu
 20 25 30
 Ser Pro Ala Gln Asn Thr Ala His Leu Asp Gln Phe Glu Arg Ile Lys
 35 40 45
 Thr Leu Gly Thr Gly Ser Phe Gly Arg Val Met Leu Val Lys His Lys
 50 55 60
 Glu Thr Gly Asn His Tyr Ala Met Lys Ile Leu Asp Lys Gln Lys Val
 65 70 75 80
 Val Lys Leu Lys Gln Ile Glu His Thr Leu Asn Glu Lys Arg Ile Leu
 85 90 95
 Gln Ala Val Asn Phe Pro Phe Leu Val Lys Leu Glu Phe Ser Phe Lys
 100 105 110
 Asp Asn Ser Asn Leu Tyr Met Val Met Glu Tyr Val Pro Gly Gly Glu
 115 120 125
 Met Phe Ser His Leu Arg Arg Ile Gly Arg Phe Arg
 130 135 140

<210> 104
 <211> 156
 <212> PRT
 <213> Homo sapiens

<400> 104

Met Val Val Phe Asn Gly Leu Leu Lys Ile Lys Ile Cys Glu Ala Val
 1 5 10 15
 Ser Leu Lys Pro Thr Ala Trp Ser Leu Arg His Ala Val Gly Pro Arg
 20 25 30

Pro Gln Thr Phe Leu Leu Asp Pro Tyr Ile Ala Leu Asn Val Asp Asp
35 40 45

Ser Arg Ile Gly Gln Thr Ala Thr Lys Gln Lys Thr Asn Ser Pro Ala
50 55 60

Trp His Asp Glu Phe Val Thr Asp Val Cys Asn Gly Arg Lys Ile Glu
65 70 75 80

Leu Ala Val Phe His Asp Ala Pro Ile Gly Tyr Asp Asp Phe Val Ala
85 90 95

Asn Cys Thr Ile Gln Phe Glu Glu Leu Leu Gln Asn Gly Ser Arg His
100 105 110

Phe Glu Asp Trp Ile Asp Leu Glu Pro Glu Gly Arg Val Tyr Val Ile
115 120 125

Ile Asp Leu Ser Gly Ser Ser Gly Glu Val Lys Ile Pro Asn Ser Ala
130 135 140

Phe Cys Glu Arg Glu Arg Val Glu Met Arg His Ser
145 150 155

<210> 105

<211> 520

<212> PRT

<213> Homo sapiens

<400> 105

Met Ile Leu Ile Pro Arg Met Leu Leu Val Leu Phe Leu Leu Leu Pro
1 5 10 15

Ile Leu Ser Ser Ala Lys Ala Gln Val Asn Pro Ala Ile Cys Arg Tyr
20 25 30

Pro Leu Gly Met Ser Gly Gly Gln Ile Pro Asp Glu Asp Ile Thr Ala
35 40 45

Ser Ser Gln Trp Ser Glu Ser Thr Ala Ala Lys Tyr Gly Arg Leu Asp
50 55 60

Ser Glu Glu Gly Asp Gly Ala Trp Cys Pro Glu Ile Pro Val Glu Pro
65 70 75 80

Asp Asp Leu Lys Glu Phe Leu Gln Ile Asp Leu His Thr Leu His Phe
85 90 95

Ile Thr Leu Val Gly Thr Gln Gly Arg His Ala Gly Gly His Gly Ile
100 105 110

Glu Phe Ala Pro Met Tyr Lys Ile Asn Tyr Ser Arg Asp Gly Thr Arg
115 120 125

Trp Ile Ser Trp Arg Asn Arg His Gly Lys Gln Val Leu Asp Gly Asn

435		440		445
Ser Asp Ser Ser Met Phe Asn Asn Asn Arg Ser Ser Ser Pro Ser Glu				
450		455		460
Gln Gly Ser Asn Ser Thr Tyr Asp Arg Ile Phe Pro Leu Arg Pro Asp				
465		470		480
Tyr Gln Glu Pro Ser Arg Leu Ile Arg Lys Leu Pro Glu Phe Ala Pro				
	485		490	495
Gly Glu Glu Glu Ser Gly Glu Asp Asp Val Val Glu Gln Gly Val Lys				
	500		505	510
Gly Glu Thr Ser Ala Ser Ile Arg				
	515		520	

<210> 106
 <211> 284
 <212> PRT
 <213> Homo sapiens

<400> 106

Met Ala Asn Phe Gln Glu His Leu Ser Cys Ser Ser Ser Pro His Leu				
1	5		10	15
Pro Phe Ser Glu Ser Lys Thr Phe Asn Gly Leu Gln Asp Glu Leu Thr				
	20		25	30
Ala Met Gly Asn His Pro Ser Pro Lys Leu Leu Glu Asp Gln Gln Glu				
	35		40	45
Lys Gly Met Val Arg Thr Glu Leu Ile Glu Ser Val His Ser Pro Val				
	50		55	60
Thr Thr Thr Val Leu Thr Ser Val Ser Glu Asp Ser Arg Asp Gln Phe				
65		70		80
Glu Asn Ser Val Leu Gln Leu Arg Glu His Asp Glu Ser Glu Thr Ala				
	85		90	95
Val Ser Gln Gly Asn Ser Asn Thr Val Asp Gly Glu Ser Thr Ser Gly				
	100		105	110
Thr Glu Asp Ile Lys Ile Gln Phe Ser Arg Ser Gly Ser Gly Ser Gly				
	115		120	125
Gly Phe Leu Glu Gly Leu Phe Gly Cys Leu Arg Pro Val Trp Asn Ile				
	130		135	140
Ile Gly Lys Ala Tyr Ser Thr Asp Tyr Lys Phe Met Gln Gln Asp Thr				
145		150		160
Trp Glu Val Pro Phe Glu Glu Ile Ser Glu Leu Gln Trp Leu Gly Ser				
	165		170	175

Gly Ala Gln Gly Ala Val Phe Leu Gly Lys Phe Arg Ala Glu Glu Val
 180 185 190
 Ala Ile Lys Lys Val Arg Glu Gln Asn Glu Thr Asp Ile Lys His Leu
 195 200 205
 Arg Lys Leu Lys His Pro Asn Ile Ile Ala Phe Lys Gly Val Cys Thr
 210 215 220
 Gln Ala Pro Cys Tyr Cys Ile Ile Met Glu Tyr Cys Ala His Gly Gln
 225 230 235 240
 Leu Tyr Glu Val Leu Arg Ala Gly Arg Lys Ile Thr Pro Arg Leu Leu
 245 250 255
 Val Asp Trp Ser Thr Gly Ile Ala Ser Gly Met Asn Tyr Leu His Leu
 260 265 270
 His Lys Ile Ile His Arg Asp Leu Lys Ser Pro Lys
 275 280

<210> 107
 <211> 185
 <212> PRT
 <213> Homo sapiens

<400> 107

Met Cys Gly Gln Arg Trp Ile His Asn Phe Thr Cys Leu Ala Phe Leu
 1 5 10 15
 Phe His Thr Leu Lys Ser Gly Asn Lys Ser Val His Leu Arg Lys Ala
 20 25 30
 Ser Ser Pro Asn Leu His Arg Arg Gln Trp Glu Lys Asn Val Pro Asn
 35 40 45
 Thr Ala Leu Thr Ala Leu Glu Asn Ala Ser Ile Leu Thr Ser Ser Leu
 50 55 60
 Thr Ala Glu Asp Asp Arg Gly Gly Ser Val Ile Lys Tyr Ser Lys Asn
 65 70 75 80
 Thr Thr Arg Lys Gln Trp Leu Lys Glu Thr Pro Asp Thr Leu Leu Asn
 85 90 95
 Ile Leu Lys Asn Ala Asp Leu Ser Leu Ala Phe Gln Thr Tyr Thr Ile
 100 105 110
 Tyr Arg Pro Gly Ser Glu Gly Phe Leu Lys Gly Pro Leu Ser Glu Glu
 115 120 125
 Thr Glu Ala Ser Asp Ser Val Asp Gly Gly His Asp Ser Val Ile Leu
 130 135 140
 Asp Pro Glu Arg Leu Glu Pro Gly Leu Asp Glu Glu Asp Thr Asp Phe
 145 150 155 160

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Glu Glu Glu Asp Asp Asn Pro Asp Trp Val Ser Glu Leu Lys Lys Arg
165 170 175

Ala Gly Trp Gln Gly Leu Cys Asp Arg
180 185

<210> 108
<211> 83
<212> PRT
<213> Homo sapiens

<400> 108

Met Ala Pro Pro Ser Glu Glu Thr Pro Leu Ile Pro Gln Arg Ser Cys
1 5 10 15

Ser Leu Leu Ser Thr Glu Ala Gly Ala Leu His Val Leu Leu Pro Ala
20 25 30

Arg Gly Pro Gly Pro Pro Gln Arg Leu Ser Phe Ser Phe Gly Val Pro
35 40 45

Val Arg Pro Val Gly Ala Asn Gly Pro Pro Leu Thr Ser Gly Phe Leu
50 55 60

Gly Gly Trp Ala Glu Ala Ser Val Gln Arg Gly Leu Trp Lys Cys Leu
65 70 75 80

Leu Thr Glu

<210> 109
<211> 213
<212> PRT
<213> Homo sapiens

<400> 109

Met Ala Glu Ser Ala Gly Ala Ser Ser Phe Phe Pro Leu Val Val Leu
1 5 10 15

Leu Leu Ala Gly Ser Gly Gly Ser Gly Pro Arg Gly Val Gln Ala Leu
20 25 30

Leu Cys Ala Cys Thr Ser Cys Leu Gln Ala Asn Tyr Thr Cys Glu Thr
35 40 45

Asp Gly Ala Cys Met Val Ser Ile Phe Asn Leu Asp Gly Met Glu His
50 55 60

His Val Arg Thr Cys Ile Pro Lys Val Glu Leu Val Pro Ala Gly Lys
65 70 75 80

Pro Phe Tyr Cys Leu Ser Ser Glu Asp Leu Arg Asn Thr His Cys Cys
85 90 95

1000
900
800
700
600
500
400
300
200
100
0

Tyr Thr Asp Tyr Cys Asn Arg Ile Asp Leu Arg Val Pro Ser Gly His
100 105 110

Leu Lys Glu Pro Glu His Pro Ser Met Trp Gly Pro Val Glu Leu Val
115 120 125

Gly Ile Ile Ala Gly Pro Val Phe Leu Leu Phe Leu Ile Ile Ile Ile
130 135 140

Val Phe Leu Val Ile Asn Tyr His Gln Arg Val Tyr His Asn Arg Gln
145 150 155 160

Arg Leu Asp Met Glu Asp Pro Ser Cys Glu Met Cys Leu Ser Lys Asp
165 170 175

Lys Thr Leu Gln Asp Leu Val Tyr Asp Leu Ser Thr Ser Gly Ser Gly
180 185 190

Ser Gly Thr Lys Phe Phe Arg Ala Ser Cys Leu Trp Leu Ala Phe Ile
195 200 205

Ser Phe Pro Ala Gly
210

<210> 110
<211> 383
<212> PRT
<213> Homo sapiens

<400> 110

Met Asp Glu Gln Glu Ala Leu Asn Ser Ile Met Asn Asp Leu Val Ala
1 5 10 15

Leu Gln Met Asn Arg Arg His Arg Met Pro Gly Tyr Glu Thr Met Lys
20 25 30

Asn Lys Asp Thr Gly His Ser Asn Arg Gln Ser Asp Val Arg Ile Lys
35 40 45

Phe Glu His Asn Gly Glu Arg Arg Ile Ile Ala Phe Ser Arg Pro Val
50 55 60

Lys Tyr Glu Asp Val Glu His Lys Val Thr Thr Val Phe Gly Gln Pro
65 70 75 80

Leu Asp Leu His Tyr Met Asn Asn Glu Leu Ser Ile Leu Leu Lys Asn
85 90 95

Gln Asp Asp Leu Asp Lys Ala Ile Asp Ile Leu Asp Arg Ser Ser Ser
100 105 110

Met Lys Ser Leu Arg Ile Leu Leu Leu Ser Gln Asp Arg Asn His Asn
115 120 125

Ser Ser Ser Pro His Ser Glu Val Ser Arg Gln Val Arg Ile Lys Ala
130 135 140

Ser Gln Ser Ala Gly Asp Ile Asn Thr Ile Tyr Gln Pro Pro Glu Pro
 145 150 155 160
 Arg Ser Arg His Leu Ser Val Ser Ser Gln Asn Pro Gly Arg Ser Ser
 165 170 175
 Pro Pro Pro Gly Tyr Val Pro Glu Arg Gln Gln His Ile Ala Arg Gln
 180 185 190
 Gly Ser Tyr Thr Ser Ile Asn Ser Glu Gly Glu Phe Ile Pro Glu Thr
 195 200 205
 Ser Glu Gln Cys Met Leu Asp Pro Leu Ser Ser Ala Glu Asn Ser Leu
 210 215 220
 Ser Gly Ser Cys Gln Ser Leu Asp Arg Ser Ala Asp Ser Pro Ser Phe
 225 230 235 240
 Arg Lys Ser Arg Met Ser Arg Ala Gln Ser Phe Pro Asp Asn Arg Gln
 245 250 255
 Glu Tyr Ser Asp Arg Glu Thr Gln Leu Tyr Asp Lys Gly Val Lys Gly
 260 265 270
 Gly Thr Tyr Pro Arg Arg Tyr His Val Ser Val His His Lys Asp Tyr
 275 280 285
 Ser Asp Gly Arg Arg Thr Phe Pro Arg Ile Arg Arg His Gln Gly Asn
 290 295 300
 Leu Phe Thr Leu Val Pro Ser Ser Arg Ser Leu Ser Thr Asn Gly Glu
 305 310 315 320
 Asn Met Gly Leu Ala Val Gln Tyr Leu Asp Pro Arg Gly Arg Leu Arg
 325 330 335
 Ser Ala Asp Ser Glu Asn Ala Leu Ser Val Gln Glu Arg Asn Val Pro
 340 345 350
 Thr Lys Cys Glu Glu Leu Ser Leu Ala Arg Arg Arg Leu Pro Arg Trp
 355 360 365
 Ser Gln Thr Ser Tyr Gly Gly Lys Gln Leu Gly Pro Trp Asp Pro
 370 375 380

<210> 111

<211> 414

<212> PRT

<213> Homo sapiens

<400> 111

Met Asp Glu Gln Glu Ala Leu Asn Ser Ile Met Asn Asp Leu Val Ala
 1 5 10 15

Leu Gln Met Asn Arg Arg His Arg Met Pro Gly Tyr Glu Thr Met Lys

20							25					30						
Asn	Lys	Asp	Thr	Gly	His	Ser	Asn	Arg	Gln	Lys	Lys	His	Asn	Ser	Ser			
35							40					45						
Ser	Ser	Ala	Leu	Leu	Asn	Ser	Pro	Thr	Val	Thr	Thr	Ser	Ser	Cys	Ala			
50							55					60						
Gly	Ala	Ser	Glu	Lys	Lys	Lys	Phe	Leu	Ser	Asp	Val	Arg	Ile	Lys	Phe			
65							70					75					80	
Glu	His	Asn	Gly	Glu	Arg	Arg	Ile	Ile	Ala	Phe	Ser	Arg	Pro	Val	Lys			
85							90					95						
Tyr	Glu	Asp	Val	Glu	His	Lys	Val	Thr	Thr	Val	Phe	Gly	Gln	Pro	Leu			
100							105					110						
Asp	Leu	His	Tyr	Met	Asn	Asn	Glu	Leu	Ser	Ile	Leu	Leu	Lys	Asn	Gln			
115							120					125						
Asp	Asp	Leu	Asp	Lys	Ala	Ile	Asp	Ile	Leu	Asp	Arg	Ser	Ser	Ser	Met			
130							135					140						
Lys	Ser	Leu	Arg	Ile	Leu	Leu	Leu	Ser	Gln	Asp	Arg	Asn	His	Asn	Ser			
145							150					155					160	
Ser	Ser	Pro	His	Ser	Glu	Val	Ser	Arg	Gln	Val	Arg	Ile	Lys	Ala	Ser			
165							170					175						
Gln	Ser	Ala	Gly	Asp	Ile	Asn	Thr	Ile	Tyr	Gln	Pro	Pro	Glu	Pro	Arg			
180							185					190						
Ser	Arg	His	Leu	Ser	Val	Ser	Ser	Gln	Asn	Pro	Gly	Arg	Ser	Ser	Pro			
195							200					205						
Pro	Pro	Gly	Tyr	Val	Pro	Glu	Arg	Gln	Gln	His	Ile	Ala	Arg	Gln	Gly			
210							215					220						
Ser	Tyr	Thr	Ser	Ile	Asn	Ser	Glu	Gly	Glu	Phe	Ile	Pro	Glu	Thr	Ser			
225							230					235					240	
Glu	Gln	Cys	Met	Leu	Asp	Pro	Leu	Ser	Ser	Ala	Glu	Asn	Ser	Leu	Ser			
245							250					255						
Gly	Ser	Cys	Gln	Ser	Leu	Asp	Arg	Ser	Ala	Asp	Ser	Pro	Ser	Phe	Arg			
260							265					270						
Lys	Ser	Arg	Met	Ser	Arg	Ala	Gln	Ser	Phe	Pro	Asp	Asn	Arg	Gln	Glu			
275							280					285						
Tyr	Ser	Asp	Arg	Glu	Thr	Gln	Leu	Tyr	Asp	Lys	Gly	Val	Lys	Gly	Gly			
290							295					300						
Thr	Tyr	Pro	Arg	Arg	Tyr	His	Val	Ser	Val	His	His	Lys	Asp	Tyr	Ser			
305							310					315					320	
Asp	Gly	Arg	Arg	Thr	Phe	Pro	Arg	Ile	Arg	Arg	His	Gln	Gly	Asn	Leu			

Gln Ile Thr Gln Gln Ser Thr Asn Gln Ser Arg Asn Pro Ala Thr Thr
180 185 190

Asn Gln Thr Glu Phe Glu Arg Val Phe
195 200

<210> 113
<211> 125
<212> PRT
<213> Homo sapiens

<400> 113

Met Ala Thr Ser Arg Tyr Glu Pro Val Ala Glu Ile Gly Val Gly Ala
1 5 10 15

Tyr Gly Thr Val Tyr Lys Ala Arg Asp Pro His Ser Gly His Phe Cys
20 25 30

Ala Leu Lys Ser Val Arg Val Pro Asn Gly Gly Gly Gly Gly Gly Gly
35 40 45

Leu Pro Ile Ser Thr Val Arg Glu Val Ala Leu Leu Arg Arg Leu Glu
50 55 60

Ala Phe Glu His Pro Asn Val Val Arg Leu Met Asp Val Cys Ala Thr
65 70 75 80

Ser Arg Thr Asp Arg Glu Ile Lys Val Thr Leu Val Phe Glu His Val
85 90 95

Asp Gln Asp Leu Arg Thr Tyr Leu Asp Lys Ala Pro Pro Pro Gly Leu
100 105 110

Pro Ala Glu Thr Ile Lys Val Ser Gly Val Gly Arg His
115 120 125

<210> 114
<211> 45
<212> PRT
<213> Homo sapiens

<400> 114

Met Ala Thr Ser Arg Tyr Glu Pro Val Ala Glu Ile Gly Val Gly Ala
1 5 10 15

Tyr Gly Thr Val Tyr Lys Ala Arg Asp Pro His Ser Gly His Phe Cys
20 25 30

Ala Leu Lys Ser Val Arg Val Pro Thr His Leu Ser Phe
35 40 45

<210> 115
<211> 160
<212> PRT
<213> Homo sapiens

<400> 115

Met Gly Val Cys Pro Gly Lys Thr Pro Phe Cys Ser Pro Lys Pro Gln
1 5 10 15

Gly Leu Ala Arg Gly His Trp Ser Arg Arg Arg Asp Ile Cys Val Thr
20 25 30

Gly Pro Leu Pro Leu Glu Pro Arg Ala Val Tyr Cys Lys Asp Val Leu
35 40 45

Asp Ile Glu Gln Phe Ser Thr Val Lys Gly Val Asn Leu Asp His Thr
50 55 60

Asp Asp Asp Phe Tyr Ser Lys Phe Ser Thr Gly Ser Val Ser Ile Pro
65 70 75 80

Trp Gln Asn Glu Met Ile Glu Thr Glu Cys Phe Lys Glu Leu Asn Val
85 90 95

Phe Gly Pro Asn Gly Thr Leu Pro Pro Asp Leu Asn Arg Asn His Pro
100 105 110

Pro Glu Pro Pro Lys Lys Gly Leu Leu Gln Arg Leu Phe Lys Arg Gln
115 120 125

His Gln Asn Asn Ser Lys Ser Ser Pro Ser Ser Lys Thr Ser Phe Asn
130 135 140

His His Ile Asn Ser Asn His Val Ser Ser Asn Ser Thr Gly Ser Ser
145 150 155 160

<210> 116

<211> 300

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(300)

<223> "XAA" can be any amino acid

<400> 116

Met Pro Arg Ala Arg Met Pro Xaa Pro Arg Ala His Ser Lys Ala Gly
1 5 10 15

Cys Pro Cys Gly Cys Pro Arg Asp Pro Leu Thr Leu Leu Ser Pro Ser
20 25 30

Gly His Ile Arg Ile Ser Asp Leu Gly Leu Ala Val Lys Ile Pro Glu
35 40 45

Gly Asp Leu Ile Arg Gly Arg Val Gly Thr Val Gly Tyr Met Ala Pro
50 55 60

Glu	Val	Leu	Asn	Asn	Gln	Arg	Tyr	Gly	Leu	Ser	Pro	Asp	Tyr	Trp	Gly	
65					70					75					80	
Leu	Gly	Cys	Leu	Ile	Tyr	Glu	Met	Ile	Glu	Gly	Gln	Ser	Pro	Phe	Arg	
				85					90					95		
Gly	Arg	Lys	Glu	Lys	Val	Lys	Arg	Glu	Glu	Val	Asp	Arg	Arg	Val	Leu	
			100					105					110			
Glu	Thr	Glu	Glu	Val	Tyr	Ser	His	Lys	Phe	Ser	Glu	Glu	Ala	Lys	Ser	
			115				120						125			
Ile	Cys	Lys	Met	Leu	Leu	Thr	Lys	Asp	Ala	Lys	Gln	Arg	Leu	Gly	Cys	
	130						135					140				
Gln	Glu	Glu	Gly	Ala	Ala	Glu	Val	Lys	Arg	His	Pro	Phe	Phe	Arg	Asn	
145					150					155					160	
Met	Asn	Phe	Lys	Arg	Leu	Glu	Ala	Gly	Met	Leu	Asp	Pro	Pro	Phe	Val	
				165					170					175		
Pro	Asp	Pro	Arg	Ala	Val	Tyr	Cys	Lys	Asp	Val	Leu	Asp	Ile	Glu	Gln	
			180					185					190			
Phe	Ser	Thr	Val	Lys	Gly	Val	Asn	Leu	Asp	His	Thr	Asp	Asp	Asp	Phe	
		195					200					205				
Tyr	Ser	Lys	Phe	Ser	Thr	Gly	Ser	Val	Ser	Ile	Pro	Trp	Gln	Asn	Glu	
	210					215					220					
Met	Ile	Glu	Thr	Glu	Cys	Phe	Lys	Glu	Leu	Asn	Val	Phe	Gly	Pro	Asn	
225					230					235					240	
Gly	Thr	Leu	Pro	Pro	Asp	Leu	Asn	Arg	Asn	His	Pro	Pro	Glu	Pro	Pro	
				245					250					255		
Lys	Lys	Gly	Leu	Leu	Gln	Arg	Leu	Phe	Lys	Arg	Gln	His	Gln	Asn	Asn	
			260					265					270			
Ser	Lys	Ser	Ser	Pro	Ser	Ser	Lys	Thr	Ser	Phe	Asn	His	His	Ile	Asn	
		275					280					285				
Ser	Asn	His	Val	Ser	Ser	Asn	Ser	Thr	Gly	Ser	Ser					
	290					295					300					

<210> 117

<211> 169

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(169)

<223> "XAA" can be any amino acid

<400> 117

Met Arg Met Pro Arg Ala Arg Met Pro Xaa Pro Arg Ala His Ser Lys
1 5 10 15
Ala Gly Cys Pro Cys Gly Cys Pro Arg Asp Pro Leu Thr Leu Leu Ser
20 25 30
Pro Ser Gly His Ile Arg Ile Ser Asp Leu Gly Leu Ala Val Lys Ile
35 40 45
Pro Glu Gly Asp Leu Ile Arg Gly Arg Val Gly Thr Val Gly Tyr Met
50 55 60
Ala Pro Glu Val Leu Asn Asn Gln Arg Tyr Gly Leu Ser Pro Asp Tyr
65 70 75 80
Trp Gly Leu Gly Cys Leu Ile Tyr Glu Met Ile Glu Gly Gln Ser Pro
85 90 95
Phe Arg Gly Arg Lys Glu Lys Val Lys Arg Glu Glu Val Asp Arg Arg
100 105 110
Val Leu Glu Thr Glu Glu Val Tyr Ser His Lys Phe Ser Glu Glu Ala
115 120 125
Lys Ser Ile Cys Lys Met Val Ser Ser Trp Trp Pro Asp Ala Thr Leu
130 135 140
Lys Leu Val Ala Pro Ser Leu Gly Leu Ala Pro Val Cys Pro Gln Asn
145 150 155 160
Ser Lys Gln Ala Glu Gly Thr Gly Val
165

<210> 118

<211> 319

<212> PRT

<213> Homo sapiens

<400> 118

Met Ala Pro Phe Leu Arg Ile Ala Phe Asn Ser Tyr Glu Leu Gly Ser
1 5 10 15
Leu Gln Ala Glu Asp Glu Ala Asn Gln Pro Phe Cys Ala Val Lys Met
20 25 30
Lys Glu Ala Leu Ser Thr Glu Arg Gly Lys Thr Leu Val Gln Lys Lys
35 40 45
Pro Thr Met Tyr Pro Glu Trp Lys Ser Thr Phe Asp Ala His Ile Tyr
50 55 60
Glu Gly Arg Val Ile Gln Ile Val Leu Met Arg Ala Ala Glu Glu Pro
65 70 75 80

Val Ser Glu Val Thr Val Gly Val Ser Val Leu Ala Glu Arg Cys Lys
85 90 95

Lys Asn Asn Gly Lys Ala Glu Phe Trp Leu Asp Leu Gln Pro Gln Ala
100 105 110

Lys Val Leu Met Ser Val Gln Tyr Phe Leu Glu Asp Val Asp Cys Lys
115 120 125

Gln Ser Met Arg Ser Glu Asp Glu Ala Lys Phe Pro Thr Met Asn Arg
130 135 140

Arg Gly Ala Ile Lys Gln Ala Lys Ile His Tyr Ile Lys Asn His Glu
145 150 155 160

Phe Ile Ala Thr Phe Phe Gly Gln Pro Thr Phe Cys Ser Val Cys Lys
165 170 175

Asp Phe Val Trp Gly Leu Asn Lys Gln Gly Tyr Lys Cys Arg Gln Cys
180 185 190

Asn Ala Ala Ile His Lys Lys Cys Ile Asp Lys Ile Ile Gly Arg Cys
195 200 205

Thr Gly Thr Ala Ala Asn Ser Arg Asp Thr Ile Phe Gln Lys Glu Arg
210 215 220

Phe Asn Ile Asp Met Pro His Arg Phe Lys Val His Asn Tyr Met Ser
225 230 235 240

Pro Thr Phe Cys Asp His Cys Gly Ser Leu Leu Leu Pro Ala Pro His
245 250 255

Asp Lys His Gln Trp Asp Cys Gly Asp Phe Cys Cys Trp Pro Arg Pro
260 265 270

Cys Pro Gln Ser Val Leu Gly Cys Arg Leu Ala Gly Leu Ser Trp Tyr
275 280 285

Phe Leu Cys Glu Leu Cys Val Asn Leu Leu Phe Leu Cys Leu Arg Arg
290 295 300

Glu Ile Val Asn Pro Val Phe His Tyr Leu Asn Val Val Ile Tyr
305 310 315

<210> 119

<211> 236

<212> PRT

<213> Homo sapiens

<400> 119

Met Asp Glu Thr His Pro Gly Tyr Gly Lys Glu Val Asp Leu Glu Phe
1 5 10 15

Leu Val Ser Pro Ser Leu Pro Cys Leu Leu Ser Phe Ala Gly Ser Ala
20 25 30

Arg His Leu Val Pro Pro Asp Ser Asn Leu Phe Ser Lys Leu Trp Ala
35 40 45

Cys Gly Val Ile Leu Phe Thr Leu Leu Ala Gly Ser Pro Pro Phe Trp
50 55 60

His Arg Arg Gln Ile Leu Met Leu Arg Met Ile Met Glu Gly Gln Tyr
65 70 75 80

Gln Phe Ser Ser Pro Glu Trp Asp Asp Arg Ser Ser Thr Val Lys Asp
85 90 95

Leu Ile Ser Arg Leu Leu Gln Val Asp Pro Glu Ala Arg Leu Thr Ala
100 105 110

Glu Gln Ala Leu Gln His Pro Phe Phe Glu Arg Cys Glu Gly Ser Gln
115 120 125

Pro Trp Asn Leu Thr Pro Arg Gln Arg Phe Arg Val Ala Val Trp Thr
130 135 140

Val Leu Ala Ala Gly Arg Val Ala Leu Ser Thr His Arg Val Arg Pro
145 150 155 160

Leu Thr Lys Asn Ala Leu Leu Arg Asp Pro Tyr Ala Leu Arg Ser Val
165 170 175

Arg His Leu Ile Asp Asn Cys Ala Phe Arg Leu Tyr Gly His Trp Val
180 185 190

Lys Lys Gly Glu Gln Gln Asn Arg Ala Ala Leu Phe Gln His Arg Pro
195 200 205

Pro Gly Pro Phe Pro Ile Met Gly Pro Glu Glu Glu Gly Asp Ser Ala
210 215 220

Ala Ile Thr Glu Asp Glu Ala Val Leu Val Leu Gly
225 230 235

<210> 120

<211> 572

<212> PRT

<213> Homo sapiens

<400> 120

Met Ala Phe Cys Ala Lys Met Arg Ser Ser Lys Lys Thr Glu Val Asn
1 5 10 15

Leu Glu Ala Pro Glu Pro Gly Val Glu Val Ile Phe Tyr Leu Ser Asp
20 25 30

Arg Glu Pro Leu Arg Leu Gly Ser Gly Glu Tyr Thr Ala Glu Glu Leu
35 40 45

Cys Ile Arg Ala Ala Gln Ala Cys Arg Ile Ser Pro Leu Cys His Asn

355					360					365						
Val	Val	Ser	Ile	Asn	Lys	Gln	Asp	Asn	Lys	Lys	Met	Glu	Leu	Lys	Leu	
370					375					380						
Ser	Ser	His	Glu	Glu	Ala	Leu	Ser	Phe	Val	Ser	Leu	Val	Asp	Gly	Tyr	
385					390					395					400	
Phe	Arg	Leu	Thr	Ala	Asp	Ala	His	His	Tyr	Leu	Cys	Thr	Asp	Val	Ala	
405					410					415						
Pro	Pro	Leu	Ile	Val	His	Asn	Ile	Gln	Asn	Gly	Cys	His	Gly	Pro	Ile	
420					425					430						
Cys	Thr	Glu	Tyr	Ala	Ile	Asn	Lys	Leu	Arg	Gln	Glu	Gly	Ser	Glu	Glu	
435					440					445						
Gly	Met	Tyr	Val	Leu	Arg	Trp	Ser	Cys	Thr	Asp	Phe	Asp	Asn	Ile	Leu	
450					455					460						
Met	Thr	Val	Thr	Cys	Phe	Glu	Lys	Ser	Glu	Gln	Val	Gln	Gly	Ala	Gln	
465					470					475					480	
Lys	Gln	Phe	Lys	Asn	Phe	Gln	Ile	Glu	Val	Gln	Lys	Gly	Arg	Tyr	Ser	
485					490					495						
Leu	His	Gly	Ser	Asp	Arg	Ser	Phe	Pro	Ser	Leu	Gly	Asp	Leu	Met	Ser	
500					505					510						
His	Leu	Lys	Lys	Gln	Ile	Leu	Arg	Thr	Asp	Asn	Ile	Ser	Phe	Met	Leu	
515					520					525						
Lys	Arg	Cys	Cys	Gln	Pro	Lys	Pro	Arg	Gly	Ser	Leu	Pro	Val	Pro	Glu	
530					535					540						
Pro	Gly	Cys	Ile	Pro	Ser	Val	Ile	Ala	Glu	Thr	Gln	Ile	Asp	Gln	Asn	
545					550					555					560	
Thr	Leu	Thr	Asp	Leu	Asn	Lys	Val	Asp	Pro	Pro	Pro					
565					570											

<210> 121
 <211> 311
 <212> PRT
 <213> Homo sapiens

<400> 121

Met	Gly	Cys	Val	Gln	Cys	Lys	Asp	Lys	Glu	Ala	Thr	Lys	Leu	Thr	Glu
1				5					10					15	
Glu	Arg	Asp	Gly	Ser	Leu	Asn	Gln	Ser	Ser	Gly	Tyr	Arg	Tyr	Gly	Thr
20					25					30					
Asp	Pro	Thr	Pro	Gln	His	Tyr	Pro	Ser	Phe	Gly	Val	Thr	Ser	Ile	Pro
35					40					45					

Asn Tyr Asn Asn Phe His Ala Ala Gly Gly Gln Gly Leu Thr Val Phe
 50 55 60
 Gly Gly Val Asn Ser Ser Ser His Thr Gly Thr Leu Arg Thr Arg Gly
 65 70 75 80
 Gly Thr Gly Val Thr Leu Phe Val Ala Leu Tyr Asp Tyr Glu Ala Arg
 85 90 95
 Thr Glu Asp Asp Leu Ser Phe His Lys Gly Glu Lys Phe Gln Ile Leu
 100 105 110
 Asn Ser Ser Glu Gly Asp Trp Trp Glu Ala Arg Ser Leu Thr Thr Gly
 115 120 125
 Glu Thr Gly Tyr Ile Pro Ser Asn Tyr Val Ala Pro Val Asp Ser Ile
 130 135 140
 Gln Ala Glu Glu Trp Tyr Phe Gly Lys Leu Gly Arg Lys Asp Ala Glu
 145 150 155 160
 Arg Gln Leu Leu Ser Phe Gly Asn Pro Arg Gly Thr Phe Leu Ile Arg
 165 170 175
 Glu Ser Glu Thr Thr Lys Gly Ala Tyr Ser Leu Ser Ile Arg Asp Trp
 180 185 190
 Asp Asp Met Lys Gly Asp His Val Lys His Tyr Lys Ile Arg Lys Leu
 195 200 205
 Asp Asn Gly Gly Tyr Tyr Ile Thr Thr Arg Ala Gln Phe Glu Thr Leu
 210 215 220
 Gln Gln Leu Val Gln His Tyr Ser Glu Arg Ala Ala Gly Leu Cys Cys
 225 230 235 240
 Arg Leu Val Val Pro Cys His Lys Gly Met Pro Arg Leu Thr Asp Leu
 245 250 255
 Ser Val Lys Thr Lys Asp Val Trp Glu Ile Pro Arg Glu Ser Leu Gln
 260 265 270
 Leu Ile Lys Arg Leu Gly Asn Gly Gln Phe Gly Glu Val Trp Met Gly
 275 280 285
 Met Leu Arg Leu Asn Tyr Ser Leu Ile Ser Phe Pro Val Trp Lys Ile
 290 295 300
 Pro Asn Thr Lys Asp Gly Arg
 305 310

<210> 122

<211> 387

<212> PRT

<213> Homo sapiens

<400> 122

Thr Met Ser Pro Glu Ser Phe Leu Glu Glu Ala Gln Ile Met Lys Lys
305 310 315 320

Leu Lys His Asp Lys Leu Val Gln Leu Tyr Ala Val Val Ser Glu Glu
325 330 335

Pro Ile Tyr Ile Val Thr Glu Tyr Met Asn Lys Gly Trp Ala Thr Pro
340 345 350

Leu Leu Ser Pro Ala His Ser Ala Leu Arg Gly Cys Leu Gly Glu Arg
355 360 365

Asn Gly Ser Phe Leu Leu Ala Thr Phe Leu Val Ser Ala Trp Val Lys
370 375 380

Tyr Ser His
385

<210> 123
<211> 516
<212> PRT
<213> Homo sapiens

<400> 123

Met Arg Leu Glu Leu Pro Ala Gly His Trp Glu Arg Pro Asp Leu Glu
1 5 10 15

Leu Leu Glu Lys Ser Thr Gln Gln Gly Arg Ala Trp Asp Leu Glu Leu
20 25 30

Leu Glu Lys Gly Ala Gly Ser Leu Pro Leu Tyr Val Trp Lys Val Ser
35 40 45

Leu Ser Leu Leu Glu Leu His Lys Arg Arg Lys Ala Leu Thr Glu Pro
50 55 60

Glu Ala Arg Tyr Tyr Leu Arg Gln Ile Val Leu Gly Cys Gln Tyr Leu
65 70 75 80

His Arg Asn Arg Val Ile His Arg Asp Leu Lys Leu Gly Asn Leu Phe
85 90 95

Leu Asn Glu Asp Leu Glu Val Lys Ile Gly Asp Phe Gly Leu Ala Thr
100 105 110

Lys Val Glu Tyr Asp Gly Glu Arg Lys Lys Thr Leu Cys Gly Thr Pro
115 120 125

Asn Tyr Ile Ala Pro Glu Val Leu Ser Lys Lys Gly His Ser Phe Glu
130 135 140

Val Asp Val Trp Ser Ile Gly Cys Ile Met Tyr Thr Leu Leu Val Gly
145 150 155 160

Lys Pro Pro Phe Glu Thr Ser Cys Leu Lys Glu Thr Tyr Leu Arg Ile

165 170 175

165								170				175			
Lys	Lys	Asn	Glu	Tyr	Ser	Ile	Pro	Lys	His	Ile	Asn	Pro	Val	Ala	Ala
180								185				190			
Ser	Leu	Ile	Gln	Lys	Met	Leu	Gln	Thr	Asp	Pro	Thr	Ala	Arg	Pro	Thr
195								200				205			
Ile	Asn	Glu	Leu	Leu	Asn	Asp	Glu	Phe	Phe	Thr	Ser	Gly	Tyr	Ile	Pro
210								215				220			
Ala	Arg	Leu	Pro	Ile	Thr	Cys	Leu	Thr	Ile	Pro	Pro	Arg	Phe	Ser	Ile
225								230				235			
Ala	Pro	Ser	Ser	Leu	Asp	Pro	Ser	Asn	Arg	Lys	Pro	Leu	Thr	Val	Leu
245								250				255			
Asn	Lys	Gly	Leu	Glu	Asn	Pro	Leu	Pro	Glu	Arg	Pro	Arg	Glu	Lys	Glu
260								265				270			
Glu	Pro	Val	Val	Arg	Glu	Thr	Gly	Glu	Val	Val	Asp	Cys	His	Leu	Ser
275								280				285			
Asp	Met	Leu	Gln	Gln	Leu	His	Ser	Val	Asn	Ala	Ser	Lys	Pro	Ser	Glu
290								295				300			
Arg	Gly	Leu	Val	Arg	Gln	Glu	Glu	Ala	Glu	Asp	Pro	Ala	Cys	Ile	Pro
305								310				315			
Ile	Phe	Trp	Val	Ser	Lys	Trp	Val	Asp	Tyr	Ser	Asp	Lys	Tyr	Gly	Leu
325								330				335			
Gly	Tyr	Gln	Leu	Cys	Asp	Asn	Ser	Val	Gly	Val	Leu	Phe	Asn	Asp	Ser
340								345				350			
Thr	Arg	Leu	Ile	Leu	Tyr	Asn	Asp	Gly	Asp	Ser	Leu	Gln	Tyr	Ile	Glu
355								360				365			
Arg	Asp	Gly	Thr	Glu	Ser	Tyr	Leu	Thr	Val	Ser	Ser	His	Pro	Asn	Ser
370								375				380			
Leu	Met	Lys	Lys	Ile	Thr	Leu	Leu	Lys	Tyr	Phe	Arg	Asn	Tyr	Met	Ser
385								390				395			
Glu	His	Leu	Leu	Lys	Ala	Gly	Ala	Asn	Ile	Thr	Pro	Arg	Glu	Gly	Asp
405								410				415			
Glu	Leu	Ala	Arg	Leu	Pro	Tyr	Leu	Arg	Thr	Trp	Phe	Arg	Thr	Arg	Ser
420								425				430			
Ala	Ile	Ile	Leu	His	Leu	Ser	Asn	Gly	Ser	Val	Gln	Ile	Asn	Phe	Phe
435								440				445			
Gln	Asp	His	Thr	Lys	Leu	Ile	Leu	Cys	Pro	Leu	Met	Ala	Ala	Val	Thr
450								455				460			
Tyr	Ile	Asp	Glu	Lys	Arg	Asp	Phe	Arg	Thr	Tyr	Arg	Leu	Ser	Leu	Leu

<210> 125
 <211> 134
 <212> PRT
 <213> Homo sapiens

<400> 125

Met	Ala	Leu	Leu	Pro	Pro	Phe	Leu	Ala	Ser	His	Arg	Leu	Glu	Val	Ser
1				5					10					15	
Arg	Asp	Ser	Gly	Trp	Leu	Gly	Gln	Cys	Trp	Leu	Gln	Gly	Val	Trp	Glu
			20					25					30		
Arg	Gly	Leu	Thr	Val	Ala	Phe	Ser	Ile	Leu	Cys	Asn	Thr	Leu	Gln	Pro
		35					40					45			
Glu	Phe	Ser	Thr	Tyr	Leu	Asn	Phe	Cys	Arg	Ser	Leu	Arg	Phe	Asp	Asp
	50					55					60				
Lys	Pro	Asp	Tyr	Ser	Tyr	Leu	Arg	Gln	Leu	Phe	Arg	Asn	Leu	Phe	His
65					70					75					80
Arg	Gln	Gly	Phe	Ser	Tyr	Asp	Tyr	Val	Phe	Asp	Trp	Asn	Met	Leu	Lys
				85					90					95	
Phe	Gly	Ala	Ser	Ser	Ser	Gln	Ala	Gln	Pro	Arg	Asp	Ser	Pro	Met	Thr
			100					105					110		
Ala	Lys	Gly	Pro	Phe	Cys	Pro	Arg	Pro	Cys	Pro	Cys	Ala	Gly	Pro	Thr
		115					120					125			
Tyr	Ser	Pro	Thr	Tyr	Trp										
	130														

<210> 126
 <211> 233
 <212> PRT
 <213> Homo sapiens

<400> 126

Met	Ala	Leu	Leu	Pro	Pro	Phe	Leu	Ala	Ser	His	Arg	Leu	Glu	Val	Ser
1				5					10					15	
Arg	Asp	Ser	Gly	Trp	Leu	Gly	Gln	Cys	Trp	Leu	Gln	Gly	Val	Trp	Glu
			20					25					30		
Arg	Gly	Leu	Thr	Val	Ala	Phe	Ser	Ile	Leu	Cys	Asn	Thr	Leu	Gln	Pro
		35					40					45			
Glu	Phe	Ser	Thr	Tyr	Leu	Asn	Phe	Cys	Arg	Ser	Leu	Arg	Phe	Asp	Asp
	50					55					60				
Lys	Pro	Asp	Tyr	Ser	Tyr	Leu	Arg	Gln	Leu	Phe	Arg	Asn	Leu	Phe	His
65					70					75					80
Arg	Gln	Gly	Phe	Ser	Tyr	Asp	Tyr	Val	Phe	Asp	Trp	Asn	Met	Leu	Lys

Ala Val Ile Val Gly Lys Glu Ser Lys Ser Ala Ala Thr Lys Glu Glu
115 120 125

Ser Val Ser Leu Lys Glu Lys Thr Lys Pro Leu Thr Pro Ser Ile Gly
130 135 140

Ala Lys Glu Lys Glu Gln His Val Ala Leu Val Thr Ser Thr Leu Pro
145 150 155 160

Pro Leu Pro Leu Pro Pro Met Leu Pro Glu Asp Lys Glu Ala Asp Ser
165 170 175

Leu Arg Gly Asn Ile Ser Val Lys Ala Val Lys Lys Glu Val Glu Lys
180 185 190

Lys Leu Arg Cys Leu Leu Ala Asp Leu Pro Leu Pro Pro Glu Leu Pro
195 200 205

Gly Gly Asp Asp Leu Ser Lys Ser Pro Glu Glu Lys Lys Thr Ala Thr
210 215 220

Gln Leu His Ser Lys Arg Arg Pro Lys Tyr Val Leu Ala Phe Tyr Leu
225 230 235 240

Leu Leu Asn

<210> 128

<211> 330

<212> PRT

<213> Homo sapiens

<400> 128

Met Ser Ala Lys Val Arg Leu Lys Lys Leu Glu Gln Leu Leu Leu Asp
1 5 10 15

Gly Pro Trp Arg Asn Glu Ser Ala Leu Ser Val Glu Thr Leu Leu Asp
20 25 30

Val Leu Val Cys Leu Tyr Thr Glu Cys Ser His Ser Ala Leu Arg Arg
35 40 45

Asp Lys Tyr Val Ala Glu Phe Leu Glu Trp Ala Lys Pro Phe Thr Gln
50 55 60

Leu Val Lys Glu Met Gln Leu His Arg Glu Asp Phe Glu Ile Ile Lys
65 70 75 80

Val Ile Gly Arg Gly Ala Phe Gly Glu Val Ala Val Val Lys Met Lys
85 90 95

Asn Thr Glu Arg Ile Tyr Ala Met Lys Ile Leu Asn Lys Trp Glu Met
100 105 110

Leu Lys Arg Ala Glu Thr Ala Cys Phe Arg Glu Glu Arg Asp Val Leu
115 120 125

115 120 125 130 135 140 145 150 155 160 165 170 175 180 185 190 195 200 205 210 215 220 225 230 235 240

Val Asn Gly Asp Cys Gln Trp Ile Thr Ala Leu His Tyr Ala Phe Gln
130 135 140

Asp Glu Asn His Leu Tyr Leu Val Met Asp Tyr Tyr Val Gly Gly Asp
145 150 155 160

Leu Leu Thr Leu Leu Ser Lys Phe Glu Asp Lys Leu Pro Glu Asp Met
165 170 175

Ala Arg Phe Tyr Ile Gly Glu Met Val Leu Ala Ile Asp Ser Ile His
180 185 190

Gln Leu His Tyr Val His Arg Asp Ile Lys Pro Asp Asn Val Leu Leu
195 200 205

Asp Val Asn Gly His Ile Arg Leu Ala Asp Phe Gly Ser Cys Leu Lys
210 215 220

Met Asn Asp Asp Gly Thr Val Gln Ser Ser Val Ala Val Gly Thr Pro
225 230 235 240

Asp Tyr Ile Ser Pro Glu Ile Leu Gln Ala Met Glu Asp Gly Met Gly
245 250 255

Lys Tyr Gly Pro Glu Cys Asp Trp Trp Ser Leu Gly Val Cys Met Tyr
260 265 270

Glu Met Leu Tyr Gly Glu Thr Pro Phe Tyr Ala Glu Ser Leu Val Glu
275 280 285

Thr Tyr Gly Lys Ile Met Asn His Glu Glu Arg Phe Gln Phe Pro Ser
290 295 300

His Val Thr Asp Val Ser Glu Glu Ala Lys Asp Leu Ile Gln Arg Leu
305 310 315 320

Ser Cys Ile Gln Arg Thr Pro Tyr Leu Gln
325 330

<210> 129

<211> 246

<212> PRT

<213> Homo sapiens

<400> 129

Met Ser Ala Lys Val Arg Leu Lys Lys Leu Glu Gln Leu Leu Leu Asp
1 5 10 15

Gly Pro Trp Arg Asn Glu Ser Ala Leu Ser Val Glu Thr Leu Leu Asp
20 25 30

Val Leu Val Cys Leu Tyr Thr Glu Cys Ser His Ser Ala Leu Arg Arg
35 40 45

Asp Lys Tyr Val Ala Glu Phe Leu Glu Trp Ala Lys Pro Phe Thr Gln

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Leu Lys Thr Val Leu Leu Leu Ala Asp Gln Met Ile Ser Arg Ile Glu
 65 70 75 80
 Tyr Ile His Ser Lys Asn Phe Ile His Arg Asp Val Lys Pro Asp Asn
 85 90 95
 Phe Leu Met Gly Leu Gly Lys Lys Gly Asn Leu Val Tyr Ile Ile Asp
 100 105 110
 Phe Gly Leu Ala Lys Lys Tyr Arg Asp Ala Arg Thr His Gln His Ile
 115 120 125
 Pro Tyr Arg Glu Asn Lys Asn Leu Thr Gly Thr Ala Arg Tyr Ala Ser
 130 135 140
 Ile Asn Thr His Leu Gly Ile Glu Gln Ser Arg Arg Asp Asp Leu Glu
 145 150 155 160
 Ser Leu Gly Tyr Val Leu Met Tyr Phe Asn Leu Gly Ser Leu Pro Trp
 165 170 175
 Gln Gly Leu Lys Ala Ala Thr Lys Arg Gln Lys Tyr Glu Arg Ile Ser
 180 185 190
 Glu Lys Lys Met Ser Thr Pro Ile Glu Val Leu Cys Lys Gly Tyr Pro
 195 200 205
 Ser Glu Phe Ala Thr Tyr Leu Asn Phe Cys Arg Ser Leu Arg Phe Asp
 210 215 220
 Asp Lys Pro Asp Tyr Ser Tyr Leu Arg Gln Leu Phe Arg Asn Leu Phe
 225 230 235 240
 His Arg Gln Gly Phe Ser Tyr Asp Tyr Val Phe Asp Trp Asn Met Leu
 245 250 255
 Lys Phe Gly Ala Ser Arg Ala Ala Asp Asp Ala Glu Arg Asp Ala Gly
 260 265 270
 Asp Arg Glu Glu Arg Leu Arg His Ser Arg Asn Pro Ala Thr Arg Gly
 275 280 285
 Leu Pro Ser Thr Ala Ser Gly Arg Leu Arg Gly Arg Arg Lys Val Ala
 290 295 300
 Pro Pro Thr Pro Leu Thr Pro Thr Ser His Thr Ala Asn Thr Ser Pro
 305 310 315 320
 Arg Pro Val Ser Gly Met Glu Arg Glu Arg Lys Val Ser Met Arg Leu
 325 330 335
 His Arg Gly Ala Pro Val Asn Ile Ser Ser Ser Asp Leu Thr Gly Arg
 340 345 350
 Gln Asp Thr Ser Arg Met Ser Thr Ser Gln Ile Pro Gly Arg Val Ala
 355 360 365

Val Met Leu Asp Leu Ala Lys Arg Ser Arg Ser Gly Lys Phe Arg Leu
 260 265 270
 Val Thr Lys Phe Lys Lys Glu Lys Asn Asn Lys Asn Lys Glu Ala His
 275 280 285
 Ser Ser Leu Gly Ala Pro Val His Leu Trp Gly Thr Glu Glu Val Ala
 290 295 300
 Ala Trp Leu Glu His Leu Ser Leu Cys Glu Tyr Lys Asp Ile Phe Thr
 305 310 315 320
 Arg His Asp Ile Arg Gly Ser Glu Leu Leu His Leu Glu Arg Arg Asp
 325 330 335
 Leu Lys Asp Leu Gly Val Thr Lys Val Gly His Met Lys Arg Ile Leu
 340 345 350
 Cys Gly Ile Lys Glu Leu Ser Arg Ser Ala Pro Ala Val Glu Ala Gln
 355 360 365
 Pro Leu Ser Ser Gln Pro Val Ala Ser Thr Ser Pro Pro Pro Arg Pro
 370 375 380
 Ser Leu Arg Pro Leu Ser Leu Trp Pro Leu Arg Leu Leu Pro Leu Arg
 385 390 395 400
 Pro Trp Ala Asp Ala Ala Ala Arg Pro Leu Leu Met Val Leu Leu Pro
 405 410 415
 Leu Ser Ala Thr Glu Ser Leu Arg Asp Thr Val His Gln Ser Ser Gly
 420 425 430
 Val Ser Asn Ile Thr Thr Gln Leu Pro Leu Lys Gln His Phe Leu Gln
 435 440 445
 Leu Arg Val Thr Trp Gly Thr Cys Val Thr Ala Thr Gln Leu Ser Pro
 450 455 460
 Ala Cys Ala Val Gly Gln Gly Ile Gln Arg Arg Leu Ala Ser Trp Ala
 465 470 475 480
 Leu Leu Ala Trp Pro Arg Ala Trp Ile Val Pro Gly Ala Pro Leu Arg
 485 490 495
 Val Ser Phe Cys Gly Arg Thr Val Trp Leu Arg Leu Leu Ala Pro Ser
 500 505 510
 Gln Phe Ser Glu Thr Trp Leu Gly Pro Ser Thr Ala Ala Cys Lys Gly
 515 520 525
 Pro Cys Leu Leu Met Gln Leu Leu Leu Asn Lys Asn Arg Ala Leu Ser
 530 535 540
 Trp Phe Glu Ser Ser Met Asp Val Ser Ser Leu Val Asp Cys Asn Leu
 545 550 555 560

Thr

<210> 132
<211> 213
<212> PRT
<213> Homo sapiens

<400> 132

Met Ser Asp Val Ala Ile Val Lys Glu Gly Trp Leu His Lys Arg Gly
1 5 10 15

Glu Tyr Ile Lys Thr Trp Arg Pro Arg Tyr Phe Leu Leu Lys Asn Asp
20 25 30

Gly Thr Phe Ile Gly Tyr Lys Glu Arg Pro Gln Asp Val Asp Gln Arg
35 40 45

Glu Ala Pro Leu Asn Asn Phe Ser Val Ala Gln Cys Gln Leu Met Lys
50 55 60

Thr Glu Arg Pro Arg Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp
65 70 75 80

Thr Thr Val Ile Glu Arg Thr Phe His Val Glu Thr Pro Glu Glu Arg
85 90 95

Glu Glu Trp Thr Thr Ala Ile Gln Thr Val Ala Asp Gly Leu Lys Lys
100 105 110

Gln Glu Glu Glu Glu Met Asp Phe Arg Ser Gly Ser Pro Ser Asp Asn
115 120 125

Ser Gly Ala Glu Glu Met Glu Val Ser Leu Ala Lys Pro Lys His Arg
130 135 140

Val Ala Leu Gly Gly Arg Ala Gly Pro Ala His Val Ser Pro His Ser
145 150 155 160

Val Ser Gln Pro Pro Trp Ala Val Cys His Gln Leu Ser Val Ile Ser
165 170 175

Leu Gly Pro Trp Ala Ser Val Gln Pro Gly Gly Thr Arg Cys Asn Leu
180 185 190

Thr Met Val Cys Trp Pro Ala Pro Ser Pro Gly Gly Gly Arg His Thr
195 200 205

Ala Ala Pro Gln His
210

<210> 133
<211> 425
<212> PRT
<213> Homo sapiens

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<400> 133

Met Ile Val His Asp Asp Val Glu Ser Glu Pro Ala Met Thr Pro Ser
1 5 10 15

Lys Glu Gly Thr Leu Ile Val Arg Gln Thr Gln Ser Ala Ser Ser Thr
20 25 30

Leu Gln Lys His Lys Ser Ser Ser Ser Phe Thr Pro Phe Ile Asp Pro
35 40 45

Arg Leu Leu Gln Ile Ser Pro Ser Ser Gly Thr Thr Val Thr Ser Val
50 55 60

Val Gly Phe Ser Cys Asp Gly Met Arg Pro Glu Ala Ile Arg Gln Asp
65 70 75 80

Pro Thr Arg Lys Gly Ser Val Val Asn Val Asn Pro Thr Asn Thr Arg
85 90 95

Pro Gln Ser Asp Thr Pro Glu Ile Arg Lys Tyr Lys Lys Arg Phe Asn
100 105 110

Ser Glu Ile Leu Cys Ala Ala Leu Trp Gly Val Asn Leu Leu Val Gly
115 120 125

Thr Glu Ser Gly Leu Met Leu Leu Asp Arg Ser Gly Gln Gly Lys Val
130 135 140

Tyr Pro Leu Ile Asn Arg Arg Arg Phe Gln Gln Met Asp Val Leu Glu
145 150 155 160

Gly Leu Asn Val Leu Val Thr Ile Ser Gly Lys Lys Asp Lys Leu Arg
165 170 175

Val Tyr Tyr Leu Ser Trp Leu Arg Asn Lys Ile Leu His Asn Asp Pro
180 185 190

Glu Val Glu Lys Lys Gln Gly Trp Thr Thr Val Gly Asp Leu Glu Gly
195 200 205

Cys Val His Tyr Lys Val Val Lys Tyr Glu Arg Ile Lys Phe Leu Val
210 215 220

Ile Ala Leu Lys Ser Ser Val Glu Val Tyr Ala Trp Ala Pro Lys Pro
225 230 235 240

Tyr His Lys Phe Met Ala Phe Lys Ser Phe Gly Glu Leu Val His Lys
245 250 255

Pro Leu Leu Val Asp Leu Thr Val Glu Glu Gly Gln Arg Leu Lys Val
260 265 270

Ile Tyr Gly Ser Cys Ala Gly Phe His Ala Val Asp Val Asp Ser Gly
275 280 285

Ser Val Tyr Asp Ile Tyr Leu Pro Thr His Ile Gln Cys Ser Ile Lys
290 295 300

Pro His Ala Ile Ile Ile Leu Pro Asn Thr Asp Gly Met Glu Leu Leu
305 310 315 320

Val Cys Tyr Glu Asp Glu Gly Val Tyr Val Asn Thr Tyr Gly Arg Ile
325 330 335

Thr Lys Asp Val Val Leu Gln Trp Gly Glu Met Pro Thr Ser Val Ala
340 345 350

Tyr Ile Arg Ser Asn Gln Thr Met Gly Trp Gly Glu Lys Ala Ile Glu
355 360 365

Ile Arg Ser Val Glu Thr Gly His Leu Asp Gly Val Phe Met His Lys
370 375 380

Arg Ala Gln Arg Leu Lys Phe Leu Cys Glu Arg Asn Asp Lys Val Phe
385 390 395 400

Phe Ala Ser Val Arg Ser Gly Gly Ser Ser Gln Val Tyr Phe Met Thr
405 410 415

Leu Gly Arg Thr Ser Leu Leu Ser Trp
420 425

<210> 134

<211> 515

<212> PRT

<213> Homo sapiens

<400> 134

Met Ala Ser Arg Thr Pro Arg Asn Cys Ala Val Leu Lys Gly Glu Val
1 5 10 15

Asp Leu Thr Ala Leu Ala Lys Glu Leu Arg Ala Val Glu Asp Val Arg
20 25 30

Pro Pro His Lys Val Thr Asp Tyr Ser Ser Ser Ser Glu Glu Ser Gly
35 40 45

Thr Thr Asp Glu Glu Asp Asp Asp Val Glu Gln Glu Gly Ala Asp Glu
50 55 60

Ser Thr Ser Gly Pro Glu Asp Thr Arg Ala Ala Ser Ser Leu Asn Leu
65 70 75 80

Ser Asn Gly Glu Thr Glu Ser Val Lys Thr Met Ile Val His Asp Asp
85 90 95

Val Glu Ser Glu Pro Ala Met Thr Pro Ser Lys Glu Gly Thr Leu Ile
100 105 110

Val Arg Gln Thr Gln Ser Ala Ser Ser Thr Leu Gln Lys His Lys Ser
115 120 125

134 515 PRT Homo sapiens 134

Ser Ser Ser Phe Thr Pro Phe Ile Asp Pro Arg Leu Leu Gln Ile Ser
 130 135 140

Pro Ser Ser Gly Thr Thr Val Thr Ser Val Val Gly Phe Ser Cys Asp
 145 150 155 160

Gly Met Arg Pro Glu Ala Ile Arg Gln Asp Pro Thr Arg Lys Gly Ser
 165 170 175

Val Val Asn Val Asn Pro Thr Asn Thr Arg Pro Gln Ser Asp Thr Pro
 180 185 190

Glu Ile Arg Lys Tyr Lys Lys Arg Phe Asn Ser Glu Ile Leu Cys Ala
 195 200 205

Ala Leu Trp Gly Val Asn Leu Leu Val Gly Thr Glu Ser Gly Leu Met
 210 215 220

Leu Leu Asp Arg Ser Gly Gln Gly Lys Val Tyr Pro Leu Ile Asn Arg
 225 230 235 240

Arg Arg Phe Gln Gln Met Asp Val Leu Glu Gly Leu Asn Val Leu Val
 245 250 255

Thr Ile Ser Gly Lys Lys Asp Lys Leu Arg Val Tyr Tyr Leu Ser Trp
 260 265 270

Leu Arg Asn Lys Ile Leu His Asn Asp Pro Glu Val Glu Lys Lys Gln
 275 280 285

Gly Trp Thr Thr Val Gly Asp Leu Glu Gly Cys Val His Tyr Lys Val
 290 295 300

Val Lys Tyr Glu Arg Ile Lys Phe Leu Val Ile Ala Leu Lys Ser Ser
 305 310 315 320

Val Glu Val Tyr Ala Trp Ala Pro Lys Pro Tyr His Lys Phe Met Ala
 325 330 335

Phe Lys Ser Phe Gly Glu Leu Val His Lys Pro Leu Leu Val Asp Leu
 340 345 350

Thr Val Glu Glu Gly Gln Arg Leu Lys Val Ile Tyr Gly Ser Cys Ala
 355 360 365

Gly Phe His Ala Val Asp Val Asp Ser Gly Ser Val Tyr Asp Ile Tyr
 370 375 380

Leu Pro Thr His Ile Gln Cys Ser Ile Lys Pro His Ala Ile Ile Ile
 385 390 395 400

Leu Pro Asn Thr Asp Gly Met Glu Leu Leu Val Cys Tyr Glu Asp Glu
 405 410 415

Gly Val Tyr Val Asn Thr Tyr Gly Arg Ile Thr Lys Asp Val Val Leu
 420 425 430

465

<210> 136
<211> 666
<212> PRT
<213> Homo sapiens

<220>
<221> -
<222> (1)..(666)
<223> "XAA" can be any amino acid

<400> 136

Met	Asp	Cys	Gln	Leu	Ser	Ile	Leu	Leu	Leu	Leu	Ser	Cys	Ser	Val	Leu
1				5					10					15	
Asp	Ser	Phe	Gly	Glu	Leu	Ile	Pro	Gln	Pro	Ser	Asn	Glu	Val	Asn	Leu
			20					25					30		
Leu	Asp	Ser	Lys	Thr	Ile	Gln	Gly	Glu	Leu	Gly	Trp	Ile	Ser	Tyr	Pro
			35				40					45			
Ser	His	Gly	Trp	Glu	Glu	Ile	Ser	Gly	Val	Asp	Glu	His	Tyr	Thr	Pro
	50					55					60				
Ile	Arg	Thr	Tyr	Gln	Val	Cys	Asn	Val	Met	Asp	His	Ser	Gln	Asn	Asn
65					70					75					80
Trp	Leu	Arg	Thr	Asn	Trp	Val	Pro	Arg	Asn	Ser	Ala	Gln	Lys	Ile	Tyr
				85					90					95	
Val	Glu	Leu	Lys	Phe	Thr	Leu	Arg	Asp	Cys	Asn	Ser	Ile	Pro	Leu	Val
			100					105					110		
Leu	Gly	Thr	Cys	Lys	Glu	Thr	Phe	Asn	Leu	Tyr	Tyr	Met	Glu	Ser	Asp
			115				120					125			
Asp	Asp	His	Gly	Val	Lys	Phe	Arg	Glu	His	Gln	Phe	Thr	Lys	Ile	Asp
	130					135					140				
Thr	Ile	Ala	Ala	Asp	Glu	Ser	Phe	Thr	Gln	Met	Asp	Leu	Gly	Asp	Arg
145					150					155					160
Ile	Leu	Lys	Leu	Asn	Thr	Glu	Ile	Arg	Glu	Val	Gly	Pro	Val	Asn	Lys
				165					170					175	
Lys	Gly	Phe	Tyr	Leu	Ala	Phe	Gln	Asp	Val	Gly	Ala	Cys	Val	Ala	Leu
			180					185					190		
Val	Ser	Val	Arg	Val	Tyr	Phe	Lys	Lys	Cys	Pro	Phe	Thr	Val	Lys	Asn
			195				200					205			
Leu	Ala	Met	Phe	Pro	Asp	Thr	Val	Pro	Met	Asp	Ser	Gln	Ser	Leu	Val
	210					215					220				

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Glu	Val	Arg	Gly	Ser	Cys	Val	Asn	Asn	Ser	Lys	Glu	Glu	Asp	Pro	Pro	225	230	235	240
Arg	Met	Tyr	Cys	Ser	Thr	Glu	Gly	Glu	Trp	Leu	Val	Pro	Ile	Gly	Lys	245	250	255	
Cys	Ser	Cys	Asn	Ala	Gly	Tyr	Glu	Glu	Arg	Gly	Phe	Met	Cys	Gln	Ala	260	265	270	
Cys	Arg	Pro	Gly	Phe	Tyr	Lys	Ala	Leu	Asp	Gly	Asn	Met	Lys	Cys	Ala	275	280	285	
Lys	Cys	Pro	Pro	His	Ser	Ser	Thr	Gln	Glu	Asp	Gly	Ser	Met	Asn	Cys	290	295	300	
Arg	Cys	Glu	Asn	Asn	Tyr	Phe	Arg	Ala	Asp	Lys	Asp	Pro	Pro	Ser	Met	305	310	315	320
Ala	Cys	Thr	Arg	Pro	Pro	Ser	Ser	Pro	Arg	Asn	Val	Ile	Ser	Asn	Ile	325	330	335	
Asn	Glu	Thr	Ser	Val	Ile	Leu	Asp	Trp	Ser	Trp	Pro	Leu	Asp	Thr	Gly	340	345	350	
Gly	Arg	Lys	Asp	Val	Thr	Phe	Asn	Ile	Ile	Cys	Lys	Lys	Cys	Gly	Trp	355	360	365	
Asn	Ile	Lys	Gln	Cys	Glu	Pro	Cys	Ser	Pro	Asn	Val	Arg	Phe	Leu	Pro	370	375	380	
Arg	Gln	Phe	Gly	Leu	Thr	Asn	Thr	Thr	Val	Thr	Val	Thr	Asp	Leu	Leu	385	390	395	400
Ala	His	Thr	Asn	Tyr	Thr	Phe	Glu	Ile	Asp	Ala	Val	Asn	Gly	Val	Ser	405	410	415	
Glu	Leu	Ser	Ser	Pro	Pro	Arg	Gln	Phe	Ala	Ala	Val	Ser	Ile	Thr	Thr	420	425	430	
Asn	Gln	Ala	Ala	Pro	Ser	Pro	Val	Leu	Thr	Ile	Lys	Lys	Asp	Arg	Thr	435	440	445	
Ser	Arg	Asn	Ser	Ile	Ser	Leu	Ser	Trp	Gln	Glu	Pro	Glu	His	Pro	Asn	450	455	460	
Gly	Ile	Ile	Leu	Asp	Tyr	Glu	Val	Lys	Tyr	Tyr	Glu	Lys	Gln	Glu	Gln	465	470	475	480
Glu	Thr	Ser	Tyr	Thr	Ile	Leu	Arg	Ala	Arg	Gly	Thr	Asn	Val	Thr	Ile	485	490	495	
Ser	Ser	Leu	Lys	Pro	Asp	Thr	Ile	Tyr	Val	Phe	Gln	Ile	Arg	Ala	Arg	500	505	510	
Thr	Ala	Ala	Gly	Tyr	Gly	Thr	Asn	Ser	Arg	Lys	Phe	Glu	Phe	Glu	Thr	515	520	525	

Met Asn Gly His Ile Arg Leu Ala Asp Phe Gly Ser Cys Leu Lys Leu
130 135 140

Met Glu Asp Gly Thr Val Gln Ser Ser Val Ala Val Gly Thr Pro Asp
145 150 155 160

Tyr Ile Ser Pro Glu Ile Leu Gln Ala Met Glu Asp Gly Lys Gly Arg
165 170 175

Tyr Gly Pro Glu Cys Asp Trp Trp Ser Leu Gly Val Cys Met Tyr Glu
180 185 190

Met Leu Tyr Gly Glu Thr Pro Phe Tyr Ala Glu Ser Leu Val Glu Thr
195 200 205

Tyr Gly Lys Ile Met Asn His Lys Glu Arg Phe Gln Phe Pro Ala Gln
210 215 220

Val Thr Asp Val Ser Glu Asn Ala Lys Asp Leu Ile Arg Arg Leu Ile
225 230 235 240

Cys Ser Arg Glu His Arg Leu Gly Gln Asn Gly Ile Glu Asp Phe Lys
245 250 255

Lys His Pro Phe Phe Ser Gly Ile Asp Trp Asp Asn Ile Arg Asn Cys
260 265 270

Glu Ala Pro Tyr Ile Pro Glu Val Ser Ser Pro Thr Asp Thr Ser Asn
275 280 285

Phe Asp Val Asp Asp Asp Cys Leu Lys Asn Ser Glu Thr Met Pro Pro
290 295 300

Pro Thr His Thr Ala Phe Ser Gly His His Leu Pro Phe Val Gly Phe
305 310 315 320

Thr Tyr Thr Ser Ser Cys Val Leu Ser Asp Arg Ser Cys Leu Arg Val
325 330 335

Thr Ala Gly Pro Thr Ser Leu Asp Leu Asp Val Asn Val Gln Arg Thr
340 345 350

Leu Asp Asn Asn Leu Ala Thr Glu Ala Tyr Glu Arg Arg Ile Lys Arg
355 360 365

Leu Glu Gln Glu Lys Leu Glu Leu Ser Arg Lys Leu Gln Glu Ser Thr
370 375 380

Gln Thr Val Gln Ala Leu Gln Tyr Ser Thr Val Asp Gly Pro Leu Thr
385 390 395 400

Ala Ser Lys Asp Leu Glu Ile Lys Asn Leu Lys Glu Glu Ile Glu Lys
405 410 415

Leu Arg Lys Gln Val Thr Glu Ser Ser His Leu Glu Gln Gln Leu Glu
420 425 430

1000 900 800 700 600 500 400 300 200 100 0

225		230		235		240
Gln Gly Ala Gln Arg Arg Pro Ala Ala Pro Pro Gln Thr Ser Trp Arg						
		245		250		255

Val Trp Arg Pro Gly Ser
260

<210> 139
<211> 203
<212> PRT
<213> Homo sapiens

<400> 139

Met Glu Val Val Asp Pro Gln Gln Leu Gly Met Phe Thr Glu Gly Glu
1 5 10 15

Leu Met Ser Val Gly Met Asp Thr Phe Ile His Arg Ile Asp Ser Thr
20 25 30

Glu Val Ile Tyr Gln Pro Arg Arg Lys Arg Ala Lys Leu Ile Gly Lys
35 40 45

Tyr Leu Met Gly Asp Leu Leu Gly Glu Gly Ser Tyr Gly Lys Val Lys
50 55 60

Glu Val Leu Asp Ser Glu Thr Leu Cys Arg Arg Ala Val Lys Ile Leu
65 70 75 80

Lys Lys Lys Lys Leu Arg Arg Ile Pro Asn Gly Glu Ala Asn Val Lys
85 90 95

Lys Glu Ile Gln Leu Leu Arg Arg Leu Arg His Lys Asn Val Ile Gln
100 105 110

Leu Val Asp Val Leu Tyr Asn Glu Glu Lys Gln Lys Met Tyr Met Val
115 120 125

Met Glu Tyr Cys Val Cys Gly Met Gln Glu Met Leu Asp Ser Val Pro
130 135 140

Glu Lys Arg Phe Pro Val Cys Gln Ala His Gly Tyr Phe Cys Gln Leu
145 150 155 160

Ile Asp Gly Leu Glu Tyr Leu His Ser Gln Gly Ile Val His Lys Asp
165 170 175

Ile Lys Pro Gly Asn Leu Leu Leu Thr Thr Gly Gly Thr Leu Lys Ile
180 185 190

Ser Asp Leu Gly Val Ala Glu Val Gly Thr Cys
195 200

<210> 140
<211> 244
<212> PRT

<213> Homo sapiens

<400> 140

Met Asp Arg Glu Thr Thr Pro Leu Gly Leu Leu Trp Leu Ile Gln Val
1 5 10 15
Ile Pro Ser Lys Leu Leu Pro Ser Leu Gln Val Lys Asp Phe Leu Ser
20 25 30
Gln Leu Arg Ser Ser Asn Arg Arg Phe Ser Ile Pro Glu Ser Gly Gln
35 40 45
Gly Gly Thr Glu Met Asp Gly Phe Arg Arg Thr Ile Glu Asn Gln His
50 55 60
Ser Arg Asn Asp Val Met Val Ser Glu Trp Leu Asn Lys Leu Asn Leu
65 70 75 80
Glu Glu Pro Pro Ser Ser Val Pro Lys Lys Cys Pro Ser Leu Thr Lys
85 90 95
Arg Ser Arg Ala Gln Glu Glu Gln Val Pro Gln Ala Trp Thr Ala Gly
100 105 110
Thr Ser Ser Asp Ser Met Ala Gln Pro Pro Gln Thr Pro Glu Thr Ser
115 120 125
Thr Phe Arg Asn Gln Met Pro Ser Pro Thr Ser Thr Gly Thr Pro Ser
130 135 140
Pro Gly Pro Arg Gly Asn Gln Gly Ala Glu Arg Gln Gly Met Asn Trp
145 150 155 160
Ser Cys Arg Thr Pro Glu Pro Asn Pro Val Thr Gly Arg Pro Leu Val
165 170 175
Asn Ile Tyr Asn Cys Ser Gly Val Gln Val Gly Asp Asn Asn Tyr Leu
180 185 190
Thr Met Gln Gln Thr Thr Ala Leu Pro Thr Trp Gly Leu Ala Pro Ser
195 200 205
Gly Lys Gly Arg Gly Leu Gln His Pro Pro Pro Val Gly Ser Gln Glu
210 215 220
Gly Pro Lys Asp Pro Glu Ala Trp Ser Arg Pro Gln Gly Trp Tyr Asn
225 230 235 240
His Ser Gly Lys

<210> 141

<211> 222

<212> PRT

<213> Homo sapiens

<400> 141

Met Val Lys Leu Tyr Leu Tyr Gln Lys Asn Val Lys Ile Ala Ile Phe
1 5 10 15

Asp Leu Lys Ser Arg Gln Asn Phe Phe Val Tyr Phe Arg Glu Glu Gln
20 25 30

Ala Arg Glu Leu Tyr Arg Arg Leu Arg Glu Lys Pro Arg Asp Gln Arg
35 40 45

Thr Glu Gly Asp Ser Gln Glu Met Val Arg Leu Leu Leu Gln Ala Ile
50 55 60

Gln Ser Phe Glu Lys Lys Val Arg Val Ile Tyr Thr Gln Leu Ser Lys
65 70 75 80

Thr Val Val Cys Lys Gln Lys Ala Leu Glu Leu Leu Pro Lys Val Glu
85 90 95

Glu Val Val Ser Leu Met Asn Glu Asp Glu Lys Thr Val Val Arg Leu
100 105 110

Gln Glu Lys Arg Gln Lys Glu Leu Trp Asn Leu Leu Lys Ile Ala Cys
115 120 125

Ser Lys Val Arg Gly Pro Val Ser Gly Ser Pro Asp Ser Met Asn Ala
130 135 140

Ser Arg Leu Ser Gln Pro Gly Gln Leu Met Ser Gln Pro Ser Thr Ala
145 150 155 160

Ser Asn Ser Leu Pro Glu Pro Ala Lys Lys Ser Glu Glu Leu Val Ala
165 170 175

Glu Ala His Asn Leu Cys Thr Leu Leu Glu Asn Ala Ile Gln Asp Thr
180 185 190

Val Arg Glu Gln Asp Gln Ser Phe Thr Ala Leu Asp Trp Ser Trp Leu
195 200 205

Gln Thr Glu Glu Glu Glu His Ser Cys Leu Glu Gln Ala Ser
210 215 220

<210> 142

<211> 409

<212> PRT

<213> Homo sapiens

<400> 142

Met Arg Leu Thr Leu Leu Cys Cys Thr Trp Arg Glu Glu Arg Met Gly
1 5 10 15

Glu Glu Gly Ser Glu Leu Pro Val Cys Ala Ser Cys Gly Gln Arg Ile
20 25 30

Tyr Asp Gly Gln Tyr Leu Gln Ala Leu Asn Ala Asp Trp His Ala Asp
 35 40 45
 Cys Phe Arg Cys Cys Asp Cys Ser Ala Ser Leu Ser His Gln Tyr Tyr
 50 55 60
 Glu Lys Asp Gly Gln Leu Phe Cys Lys Lys Asp Tyr Trp Ala Arg Tyr
 65 70 75 80
 Gly Glu Ser Cys His Gly Cys Ser Glu Gln Ile Thr Lys Gly Leu Val
 85 90 95
 Met Val Ala Gly Glu Leu Lys Tyr His Pro Glu Cys Phe Ile Cys Leu
 100 105 110
 Thr Cys Gly Thr Phe Ile Gly Asp Gly Asp Thr Tyr Thr Leu Val Glu
 115 120 125
 His Ser Lys Leu Tyr Cys Gly His Cys Tyr Tyr Gln Thr Val Val Thr
 130 135 140
 Pro Val Ile Glu Gln Ile Leu Pro Asp Ser Pro Gly Ser His Leu Pro
 145 150 155 160
 His Thr Val Thr Leu Val Ser Ile Pro Ala Ser Ser His Gly Lys Arg
 165 170 175
 Gly Leu Ser Val Ser Ile Asp Pro Pro His Gly Pro Pro Gly Cys Gly
 180 185 190
 Thr Glu His Ser His Thr Val Arg Val Gln Gly Val Asp Pro Gly Cys
 195 200 205
 Met Ser Pro Asp Val Lys Asn Ser Ile His Val Gly Asp Arg Ile Leu
 210 215 220
 Glu Ile Asn Gly Thr Pro Ile Arg Asn Val Pro Leu Asp Glu Ile Asp
 225 230 235 240
 Leu Leu Ile Gln Glu Thr Ser Arg Leu Leu Gln Leu Thr Leu Glu His
 245 250 255
 Asp Pro His Asp Thr Leu Gly His Gly Leu Gly Pro Glu Thr Ser Pro
 260 265 270
 Leu Ser Ser Pro Ala Tyr Thr Pro Ser Gly Glu Ala Gly Ser Ser Ala
 275 280 285
 Arg Gln Lys Pro Val Leu Arg Ser Cys Ser Ile Asp Arg Ser Pro Gly
 290 295 300
 Ala Gly Ser Leu Gly Ser Pro Ala Ser Gln Arg Lys Asp Leu Gly Arg
 305 310 315 320
 Ser Glu Ser Leu Arg Val Val Cys Arg Pro His Arg Ile Phe Arg Pro
 325 330 335

Ser Asp Leu Ile His Gly Glu Val Leu Gly Lys Gly Cys Phe Gly Gln
340 345 350

Ala Ile Lys Val Gln Ser Met Pro Gly Ser Gln Leu Asp Ser Leu Gly
355 360 365

Gly Thr Pro Pro Ser Ser Phe Leu Pro Ser Leu Trp Lys His Ser Gly
370 375 380

Arg Gly Ile Trp Leu Ser Asp Ser Leu Ala Ser Ala Leu Ser Ser Leu
385 390 395 400

Gly Leu Leu Glu Leu Ile Arg Asn Arg
405

<210> 143

<211> 305

<212> PRT

<213> Homo sapiens

<400> 143

Met Arg Leu Thr Leu Leu Cys Cys Thr Trp Arg Glu Glu Arg Met Gly
1 5 10 15

Glu Glu Gly Ser Glu Leu Pro Val Cys Ala Ser Cys Gly Gln Arg Ile
20 25 30

Tyr Asp Gly Gln Tyr Leu Gln Ala Leu Asn Ala Asp Trp His Ala Asp
35 40 45

Cys Phe Arg Cys Cys Asp Cys Ser Ala Ser Leu Ser His Gln Tyr Tyr
50 55 60

Glu Lys Asp Gly Gln Leu Phe Cys Lys Lys Asp Tyr Trp Ala Arg Tyr
65 70 75 80

Gly Glu Ser Cys His Gly Cys Ser Glu Gln Ile Thr Lys Gly Leu Val
85 90 95

Met Val Ala Gly Glu Leu Lys Tyr His Pro Glu Cys Phe Ile Cys Leu
100 105 110

Thr Cys Gly Thr Phe Ile Gly Asp Gly Asp Thr Tyr Thr Leu Val Glu
115 120 125

His Ser Lys Leu Tyr Cys Gly His Cys Tyr Tyr Gln Thr Val Val Thr
130 135 140

Pro Val Ile Glu Gln Ile Leu Pro Asp Ser Pro Gly Ser His Leu Pro
145 150 155 160

His Thr Val Thr Leu Val Ser Ile Pro Ala Ser Ser His Gly Lys Arg
165 170 175

Gly Leu Ser Val Ser Ile Asp Pro Pro His Gly Pro Pro Gly Cys Gly
180 185 190

Thr Glu His Ser His Thr Val Arg Val Gln Gly Val Asp Pro Gly Cys
195 200 205

Met Ser Pro Asp Val Lys Asn Ser Ile His Val Gly Asp Arg Ile Leu
210 215 220

Glu Ile Asn Gly Thr Pro Ile Arg Asn Val Pro Leu Asp Glu Ile Asp
225 230 235 240

Leu Leu Ile Gln Glu Thr Ser Arg Leu Leu Gln Leu Thr Leu Glu His
245 250 255

Asp Pro His Asp Thr Leu Gly His Gly Leu Gly Pro Glu Thr Ser Pro
260 265 270

Leu Ser Ser Pro Ala Tyr Thr Pro Ser Gly Glu Ala Gly Ser Ser Ala
275 280 285

Arg Gln Lys Pro Val Phe Ala Arg Thr Trp Val Ala Leu Ser Pro Ser
290 295 300

Ala
305

<210> 144
<211> 780
<212> PRT
<213> Homo sapiens

<400> 144

Met Ala Ser Asp Ala Val Gln Ser Glu Pro Arg Ser Trp Ser Leu Leu
1 5 10 15

Glu Gln Leu Gly Leu Ala Gly Ala Asp Leu Ala Ala Pro Gly Val Gln
20 25 30

Gln Gln Leu Glu Leu Glu Arg Glu Arg Leu Arg Arg Glu Ile Arg Lys
35 40 45

Glu Leu Lys Leu Lys Glu Gly Ala Glu Asn Leu Arg Arg Ala Thr Thr
50 55 60

Asp Leu Gly Arg Ser Leu Gly Pro Val Glu Leu Leu Leu Arg Gly Ser
65 70 75 80

Ser Arg Arg Leu Asp Leu Leu His Gln Gln Leu Gln Glu Leu His Ala
85 90 95

His Val Val Leu Pro Asp Pro Ala Ala Thr His Asp Gly Pro Gln Ser
100 105 110

Pro Gly Ala Gly Gly Pro Thr Cys Ser Ala Thr Asn Leu Ser Arg Val
115 120 125

Ala Gly Leu Glu Lys Gln Leu Ala Ile Glu Leu Lys Val Lys Gln Gly

144
780
PRT
Homo sapiens
144

130	135	140
Ala Glu Asn Met Ile Gln Thr Tyr Ser Asn Gly Ser Thr Lys Asp Arg		
145	150	155 160
Lys Leu Leu Leu Thr Ala Gln Gln Met Leu Gln Asp Ser Lys Thr Lys		
	165 170	175
Ile Asp Ile Ile Arg Met Gln Leu Arg Arg Ala Leu Gln Ala Asp Gln		
	180 185	190
Leu Glu Asn Gln Ala Ala Pro Asp Asp Thr Gln Gly Ser Pro Asp Leu		
	195 200	205
Gly Ala Val Glu Leu Arg Ile Glu Glu Leu Arg His His Phe Arg Val		
	210 215	220
Glu His Ala Val Ala Glu Gly Ala Lys Asn Val Leu Arg Leu Leu Ser		
225	230 235	240
Ala Ala Lys Ala Pro Asp Arg Lys Ala Val Ser Glu Ala Gln Glu Lys		
	245 250	255
Leu Thr Glu Ser Asn Gln Lys Leu Gly Leu Leu Arg Glu Ala Leu Glu		
	260 265	270
Arg Arg Leu Gly Glu Leu Pro Ala Asp His Pro Lys Gly Arg Leu Leu		
	275 280	285
Arg Glu Glu Leu Ala Ala Ala Ser Ser Ala Ala Phe Ser Thr Arg Leu		
	290 295	300
Ala Gly Pro Phe Pro Ala Thr His Tyr Ser Thr Leu Cys Lys Pro Ala		
305	310 315	320
Pro Leu Thr Gly Thr Leu Glu Val Arg Val Val Gly Cys Arg Asp Leu		
	325 330	335
Pro Glu Thr Ile Pro Trp Asn Pro Thr Pro Ser Met Gly Gly Pro Gly		
	340 345	350
Thr Pro Asp Ser Arg Pro Pro Phe Leu Ser Arg Pro Ala Arg Gly Leu		
	355 360	365
Tyr Ser Arg Ser Gly Ser Leu Ser Gly Arg Ser Ser Leu Lys Ala Glu		
	370 375	380
Ala Glu Asn Thr Ser Glu Val Ser Thr Val Leu Lys Leu Asp Asn Thr		
385	390 395	400
Val Val Gly Gln Thr Ser Trp Lys Pro Cys Gly Pro Asn Ala Trp Asp		
	405 410	415
Gln Ser Phe Thr Leu Glu Leu Glu Arg Ala Arg Glu Leu Glu Leu Ala		
	420 425	430
Val Phe Trp Arg Asp Gln Arg Gly Leu Cys Ala Leu Lys Phe Leu Lys		

435					440					445					
Leu	Glu	Asp	Phe	Leu	Asp	Asn	Glu	Arg	His	Glu	Val	Gln	Leu	Asp	Met
450						455					460				
Glu	Pro	Gln	Gly	Cys	Leu	Val	Ala	Glu	Val	Thr	Phe	Arg	Asn	Pro	Val
465					470					475					480
Ile	Glu	Arg	Ile	Pro	Arg	Leu	Arg	Arg	Gln	Lys	Lys	Ile	Phe	Ser	Lys
				485					490					495	
Gln	Gln	Gly	Lys	Ala	Phe	Gln	Arg	Ala	Arg	Gln	Met	Asn	Ile	Asp	Val
			500					505						510	
Ala	Thr	Trp	Val	Arg	Leu	Leu	Arg	Arg	Leu	Ile	Pro	Asn	Ala	Thr	Gly
			515				520						525		
Thr	Gly	Thr	Phe	Ser	Pro	Gly	Ala	Ser	Pro	Gly	Ser	Glu	Ala	Arg	Thr
	530					535					540				
Thr	Gly	Asp	Ile	Ser	Val	Glu	Lys	Leu	Asn	Leu	Gly	Thr	Asp	Ser	Asp
545					550					555					560
Ser	Ser	Pro	Gln	Lys	Ser	Ser	Arg	Asp	Pro	Pro	Ser	Ser	Pro	Ser	Ser
				565					570					575	
Leu	Ser	Ser	Pro	Ile	Gln	Glu	Ser	Thr	Ala	Pro	Glu	Leu	Pro	Ser	Glu
			580					585					590		
Thr	Gln	Glu	Thr	Pro	Gly	Pro	Ala	Leu	Cys	Ser	Pro	Leu	Arg	Lys	Ser
			595				600						605		
Pro	Leu	Thr	Leu	Glu	Asp	Phe	Lys	Phe	Leu	Ala	Val	Leu	Gly	Arg	Gly
	610					615					620				
His	Phe	Gly	Lys	Val	Leu	Leu	Ser	Glu	Phe	Arg	Pro	Ser	Gly	Glu	Leu
625					630					635					640
Phe	Ala	Ile	Lys	Ala	Leu	Lys	Lys	Gly	Asp	Ile	Val	Ala	Arg	Asp	Glu
				645					650					655	
Val	Glu	Ser	Leu	Met	Cys	Glu	Lys	Arg	Ile	Leu	Ala	Ala	Val	Thr	Ser
			660					665					670		
Ala	Gly	His	Pro	Phe	Leu	Val	Asn	Leu	Phe	Gly	Cys	Phe	Gln	Thr	Pro
			675				680						685		
Glu	His	Val	Cys	Phe	Val	Met	Glu	Tyr	Ser	Ala	Gly	Gly	Asp	Leu	Met
	690					695					700				
Leu	His	Ile	His	Ser	Asp	Val	Phe	Ser	Glu	Pro	Arg	Ala	Ile	Phe	Tyr
705					710					715					720
Ser	Ala	Cys	Arg	Leu	Pro	Pro	Pro	Phe	Val	Pro	Thr	Leu	Ser	Gly	Arg
				725					730					735	
Thr	Asp	Val	Ser	Asn	Phe	Asp	Glu	Glu	Phe	Thr	Gly	Glu	Ala	Pro	Thr

	740		745		750										
Leu	Ser	Pro	Pro	Arg	Asp	Ala	Arg	Pro	Leu	Thr	Ala	Ala	Glu	Gln	Ala
		755					760					765			

Ala	Phe	Leu	Asp	Phe	Asp	Phe	Val	Ala	Gly	Gly	Cys
	770					775					780

<210> 145
 <211> 401
 <212> PRT
 <213> Homo sapiens

<400> 145

Met	Ala	Ser	Asp	Ala	Val	Gln	Ser	Glu	Pro	Arg	Ser	Trp	Ser	Leu	Leu
1				5				10						15	

Glu	Gln	Leu	Gly	Leu	Ala	Gly	Ala	Asp	Leu	Ala	Ala	Pro	Gly	Val	Gln
		20						25					30		

Gln	Gln	Leu	Glu	Leu	Glu	Arg	Glu	Arg	Leu	Arg	Arg	Glu	Ile	Arg	Lys
		35					40					45			

Glu	Leu	Lys	Leu	Lys	Glu	Gly	Ala	Glu	Asn	Leu	Arg	Arg	Ala	Thr	Thr
	50					55					60				

Asp	Leu	Gly	Arg	Ser	Leu	Gly	Pro	Val	Glu	Leu	Leu	Leu	Arg	Gly	Ser
65					70					75					80

Ser	Arg	Arg	Leu	Asp	Leu	Leu	His	Gln	Gln	Leu	Gln	Glu	Leu	His	Ala
				85					90					95	

His	Val	Val	Leu	Pro	Asp	Pro	Ala	Ala	Thr	His	Asp	Gly	Pro	Gln	Ser
			100					105					110		

Pro	Gly	Ala	Gly	Gly	Pro	Thr	Cys	Ser	Ala	Thr	Asn	Leu	Ser	Arg	Val
		115					120					125			

Ala	Gly	Leu	Glu	Lys	Gln	Leu	Ala	Ile	Glu	Leu	Lys	Val	Lys	Gln	Gly
	130					135					140				

Ala	Glu	Asn	Met	Ile	Gln	Thr	Tyr	Ser	Asn	Gly	Ser	Thr	Lys	Asp	Arg
145					150					155					160

Lys	Leu	Leu	Leu	Thr	Ala	Gln	Gln	Met	Leu	Gln	Asp	Ser	Lys	Thr	Lys
				165					170					175	

Ile	Asp	Ile	Ile	Arg	Met	Gln	Leu	Arg	Arg	Ala	Leu	Gln	Ala	Asp	Gln
			180					185					190		

Leu	Glu	Asn	Gln	Ala	Ala	Pro	Asp	Asp	Thr	Gln	Gly	Ser	Pro	Asp	Leu
		195					200					205			

Gly	Ala	Val	Glu	Leu	Arg	Ile	Glu	Glu	Leu	Arg	His	His	Phe	Arg	Val
	210					215					220				

00746404

Glu His Ala Val Ala Glu Gly Ala Lys Asn Val Leu Arg Leu Leu Ser
225 230 235 240

Ala Ala Lys Ala Pro Asp Arg Lys Ala Val Ser Glu Ala Gln Glu Lys
245 250 255

Leu Thr Glu Ser Asn Gln Lys Leu Gly Leu Leu Arg Glu Ala Leu Glu
260 265 270

Arg Arg Leu Gly Glu Leu Pro Ala Asp His Pro Lys Gly Arg Leu Leu
275 280 285

Arg Glu Glu Leu Ala Ala Ala Ser Ser Ala Ala Phe Ser Thr Arg Leu
290 295 300

Ala Gly Pro Phe Pro Ala Thr His Tyr Ser Thr Leu Cys Lys Pro Ala
305 310 315 320

Pro Leu Thr Gly Thr Leu Glu Val Arg Val Val Gly Cys Arg Asp Leu
325 330 335

Pro Glu Thr Ile Pro Trp Asn Pro Thr Pro Ser Met Gly Gly Pro Gly
340 345 350

Thr Pro Asp Ser Arg Pro Pro Phe Leu Ser Arg Pro Ala Arg Gly Leu
355 360 365

Tyr Ser Arg Ser Gly Ser Leu Ser Gly Arg Ser Ser Leu Lys Ala Glu
370 375 380

Ala Glu Asn Thr Ser Glu Val Ser Thr Val Leu Lys Leu Asp Asn Thr
385 390 395 400

His

<210> 146

<211> 96

<212> PRT

<213> Homo sapiens

<400> 146

Met Gln Ser Phe Leu Val Glu Gly Arg Phe Lys His Glu Met Phe Glu
1 5 10 15

Lys Val Phe Ala Glu Glu Arg Asn Gly Gly Gln Arg Leu Leu Cys Ala
20 25 30

Thr Asp Val Pro Ile Arg Thr Val Ser Ser Ala Ala Ser Gln Gly Leu
35 40 45

His Met Gln Asn Asp Asp Ala Cys Leu Gly Ala Ala Ser Pro Ser Ala
50 55 60

Ala Ser Trp Ser Arg Arg Ser Ala Glu Ser Lys Val Ser Leu Cys Trp
65 70 75 80

Lys Leu Lys Trp Lys Glu Asp Leu Val Trp Phe Tyr Ser Gln Ser His
85 90 95

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<210> 147
<211> 333
<212> PRT
<213> Homo sapiens
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<400> 147

Met His Arg Tyr Phe Glu Ser Pro Arg Arg Leu Leu Pro Val His Phe
1 5 10 15

Cys Cys Cys Gln Trp Arg Gly Gly Gly Val Asp Phe Glu Cys Leu Leu
20 25 30

Gly Gly Val Trp Asp Arg Cys Arg Lys Val Leu Arg Ala Gln Glu Cys
35 40 45

Glu Trp Pro Arg His Leu Pro Ser Ala Cys Leu Leu Ser Ser Ala Cys
50 55 60

Arg Gly Gln Pro Glu Arg Arg Ala Ala Val Val Gly Ala Gln Asp Pro
65 70 75 80

Thr Glu Pro Pro Arg Leu Ser Arg Ser Leu Ser Gly Ala Ser Pro Phe
85 90 95

Leu Gly Glu Thr Lys Gln Glu Thr Leu Thr Asn Ile Ser Ala Val Asn
100 105 110

Tyr Asp Phe Asp Glu Glu Tyr Phe Ser Asn Thr Ser Glu Leu Ala Lys
115 120 125

Asp Phe Ile Arg Arg Leu Leu Val Lys Asp Pro Lys Arg Arg Met Thr
130 135 140

Ile Ala Gln Ser Leu Glu His Ser Trp Ile Lys Ala Ile Arg Arg Arg
145 150 155 160

Asn Val Arg Gly Glu Asp Ser Gly Arg Lys Pro Glu Arg Arg Arg Leu
165 170 175

Lys Thr Thr Arg Leu Lys Glu Tyr Thr Ile Lys Ser His Ser Ser Leu
180 185 190

Pro Pro Asn Asn Ser Tyr Ala Asp Phe Glu Arg Phe Ser Lys Val Leu
195 200 205

Glu Glu Ala Ala Ala Ala Glu Glu Gly Leu Arg Glu Leu Gln Arg Ser
210 215 220

Arg Arg Leu Cys His Glu Asp Val Glu Ala Leu Ala Ala Ile Tyr Glu
225 230 235 240

Glu Lys Glu Ala Trp Tyr Arg Glu Glu Ser Asp Ser Leu Gly Gln Asp

[illegible]

	245		250		255
Leu Arg Arg	Leu Arg Gln Glu Leu	Leu Lys Thr Glu Ala	Leu Lys Arg		
260		265		270	
Gln Ala Gln Glu Glu Ala Lys Gly Ala Leu Leu Gly Thr Ser Gly Leu					
275		280		285	
Lys Arg Arg Phe Ser Arg Leu Glu Asn Arg Tyr Glu Ala Leu Ala Lys					
290		295		300	
Gln Val Ala Ser Glu Met Arg Phe Val Gln Asp Leu Val Arg Ala Leu					
305		310		315	320
Glu Gln Glu Lys Leu Gln Gly Val Glu Cys Gly Leu Arg					
	325		330		

<210> 148
 <211> 131
 <212> PRT
 <213> Homo sapiens

<400> 148

Met Leu Lys Glu Phe Leu Glu Ile Pro Phe Pro Thr Ser Pro Glu Cys					
1	5		10		15
Thr Leu Gln Pro Lys Ser Gln Gln Pro Thr Gly Lys Glu Ala Glu Glu					
20		25		30	
His Pro Thr Ser Ala Pro Leu Thr His Ser Leu Leu Pro Pro Thr Pro					
35		40		45	
Leu Trp Val Val Ser His Phe Ile Phe Asp Phe Arg Gly Glu Thr Ala					
50		55		60	
Leu His Lys Ala Ala Cys Gln Arg Asn Arg Ala Val Cys Gln Leu Leu					
65		70		75	80
Val Asp Ala Gly Ala Ser Leu Arg Lys Thr Asp Ser Lys Gly Lys Thr					
	85		90		95
Pro Gln Glu Arg Ala Gln Gln Ala Gly Asp Pro Asp Leu Ala Ala Tyr					
100		105		110	
Leu Glu Ser Arg Gln Asn Tyr Lys Val Ile Gly His Glu Asp Leu Glu					
115		120		125	

Thr Ala Val
 130

<210> 149
 <211> 272
 <212> PRT
 <213> Homo sapiens

<400> 149

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Met 1	Arg	Gly	Ala	Ala 5	Arg	Leu	Gly	Arg	Pro 10	Gly	Arg	Ser	Cys	Leu 15	Pro
Gly	Pro	Ala	Leu 20	Arg	Ala	Pro	Pro	Arg 25	Pro	Pro	Leu	Leu	Leu 30	Leu	Leu
Ala	Leu	Leu 35	Pro	Leu	Leu	Pro	Ala 40	Pro	Gly	Ala	Ala	Ala 45	Ala	Pro	Ala
Pro	Arg 50	Pro	Pro	Glu	Leu	Gln 55	Ser	Ala	Ser	Ala	Gly 60	Pro	Ser	Val	Ser
Leu 65	Tyr	Leu	Ser	Glu	Asp 70	Glu	Val	Arg	Arg	Leu 75	Ile	Gly	Leu	Asp	Ala 80
Glu	Leu	Tyr	Tyr	Val 85	Arg	Asn	Asp	Leu	Ile 90	Ser	His	Tyr	Ala	Leu 95	Ser
Phe	Ser	Leu	Leu 100	Val	Pro	Ser	Glu	Thr 105	Asn	Phe	Leu	His	Phe	Thr	Trp
His	Ala	Lys 115	Ser	Lys	Val	Glu	Tyr 120	Lys	Leu	Gly	Phe	Gln 125	Val	Asp	Asn
Val 130	Leu	Ala	Met	Asp	Met	Pro 135	Gln	Val	Asn	Ile	Ser 140	Val	Gln	Gly	Glu
Val 145	Pro	Arg	Thr	Leu	Ser 150	Val	Phe	Arg	Val	Glu 155	Leu	Ser	Cys	Thr	Gly 160
Lys	Val	Asp	Ser	Glu 165	Val	Met	Ile	Leu	Met 170	Gln	Leu	Asn	Leu	Thr 175	Val
Asn	Ser	Ser	Lys 180	Asn	Phe	Thr	Val	Leu 185	Asn	Phe	Lys	Arg	Arg 190	Lys	Met
Cys	Tyr	Lys 195	Lys	Leu	Glu	Glu 200	Val	Lys	Thr	Ser	Ala	Leu 205	Asp	Lys	Asn
Thr	Ser 210	Arg	Thr	Ile	Tyr	Asp 215	Pro	Val	His	Ala	Ala 220	Pro	Thr	Thr	Ser
Thr 225	Arg	Val	Phe	Tyr	Ile 230	Ser	Val	Gly	Val	Cys 235	Cys	Ala	Val	Ile	Phe 240
Leu	Val	Ala	Ile	Ile 245	Leu	Ala	Val	Leu	His 250	Leu	His	Ser	Met	Lys 255	Arg
Ile	Glu	Leu	Asp 260	Asp	Arg	Tyr	Cys	Thr 265	Tyr	Phe	Gly	Lys	Glu 270	Lys	Lys

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<210> 150
<211> 344
<212> PRT
<213> Homo sapiens
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<400> 150

Met Pro Gln Val Asn Ile Ser Val Gln Gly Glu Val Pro Arg Thr Leu
1 5 10 15

Ser Val Phe Arg Val Glu Leu Ser Cys Thr Gly Lys Val Asp Ser Glu
20 25 30

Val Met Ile Leu Met Gln Leu Asn Leu Thr Val Asn Ser Ser Lys Asn
35 40 45

Phe Thr Val Leu Asn Phe Lys Arg Arg Lys Met Cys Tyr Lys Lys Leu
50 55 60

Glu Glu Val Lys Thr Ser Ala Leu Asp Lys Asn Thr Ser Arg Thr Ile
65 70 75 80

Tyr Asp Pro Val His Ala Ala Pro Thr Thr Ser Thr Arg Val Phe Tyr
85 90 95

Ile Ser Val Gly Val Cys Cys Ala Val Ile Phe Leu Val Ala Ile Ile
100 105 110

Leu Ala Val Leu His Leu His Ser Met Lys Arg Ile Glu Leu Asp Asp
115 120 125

Ser Ile Ser Ala Ser Ser Ser Ser Gln Gly Leu Ser Gln Pro Ser Thr
130 135 140

Gln Thr Thr Gln Tyr Leu Arg Ala Asp Thr Pro Asn Asn Ala Thr Pro
145 150 155 160

Ile Thr Ser Ser Tyr Tyr Pro Thr Leu Arg Ile Glu Lys Asn Asp Leu
165 170 175

Arg Ser Val Thr Leu Leu Glu Ala Lys Gly Lys Val Lys Asp Ile Ala
180 185 190

Ile Ser Arg Glu Arg Ile Thr Leu Lys Asp Val Leu Gln Glu Gly Thr
195 200 205

Phe Gly Arg Ile Phe His Gly Ile Leu Ile Asp Glu Lys Asp Pro Asn
210 215 220

Lys Glu Lys Gln Ala Phe Val Lys Thr Val Lys Asp Gln Ala Ser Glu
225 230 235 240

Ile Gln Val Thr Met Met Leu Thr Glu Ser Cys Lys Leu Arg Gly Leu
245 250 255

His His Arg Asn Leu Leu Pro Ile Thr His Val Cys Ile Glu Glu Gly
260 265 270

Glu Lys Pro Met Val Ile Leu Pro Tyr Met Asn Trp Gly Asn Leu Lys
275 280 285

Leu Phe Leu Arg Gln Cys Lys Leu Val Glu Ala Asn Asn Pro Gln Ala

Val Ala Val Gly Gln Cys Pro Leu Val Gly Pro Gly Pro Met His Arg
 35 40 45

Arg His Leu Leu Leu Pro Ala Arg Val Pro Arg Pro Gly Leu Ser Glu
 50 55 60

Ala Leu Ser Leu Leu Leu Phe Ala Val Val Leu Ser Arg Ala Ala Ala
 65 70 75 80

Leu Gly Cys Ile Gly Leu Val Ala His Ala Gly Gln Leu Thr Ala Val
 85 90 95

Trp Arg Arg Pro Gly Ala Ala Arg Ala Pro
 100 105

<210> 153
 <211> 50
 <212> PRT
 <213> Homo sapiens

<400> 153

Met Val Asn Leu Ser His Glu Asp Phe Glu Phe Ile Ser Gly Thr Arg
 1 5 10 15

Met Arg Lys Leu Ala Arg Glu Gly Gln Lys Pro Pro Glu Gly Phe Met
 20 25 30

Ala Pro Lys Ala Trp Thr Val Leu Thr Glu Tyr Tyr Lys Ser Leu Glu
 35 40 45

Lys Ala
 50

<210> 154
 <211> 238
 <212> PRT
 <213> Homo sapiens

<400> 154

Met Ala Arg Thr Thr Ser Gln Leu Tyr Asp Ala Val Pro Ile Gln Ser
 1 5 10 15

Ser Val Val Leu Cys Ser Cys Pro Ser Pro Ser Met Val Arg Thr Gln
 20 25 30

Thr Glu Ser Ser Thr Pro Pro Gly Ile Pro Gly Gly Ser Arg Gln Gly
 35 40 45

Pro Ala Met Asp Gly Thr Ala Ala Glu Pro Arg Pro Gly Ala Gly Ser
 50 55 60

Leu Gln His Ala Gln Pro Pro Pro Gln Pro Arg Lys Lys Arg Pro Glu
 65 70 75 80

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Asp Phe Lys Phe Gly Lys Ile Leu Gly Glu Gly Ser Phe Ser Thr Val
85 90 95

Val Leu Ala Arg Glu Leu Ala Thr Ser Arg Glu Tyr Ala Ile Lys Ile
100 105 110

Leu Glu Lys Arg His Ile Ile Lys Glu Asn Lys Val Pro Tyr Val Thr
115 120 125

Arg Glu Arg Asp Val Met Ser Arg Leu Asp His Pro Phe Phe Val Lys
130 135 140

Leu Tyr Phe Thr Phe Gln Asp Asp Glu Lys Leu Tyr Phe Gly Leu Ser
145 150 155 160

Tyr Ala Lys Asn Gly Glu Leu Leu Lys Tyr Ile Arg Lys Ile Gly Ser
165 170 175

Phe Asp Glu Thr Cys Thr Arg Phe Tyr Thr Ala Glu Ile Val Ser Ala
180 185 190

Leu Glu Tyr Leu His Gly Lys Gly Ile Ile His Arg Asp Leu Lys Pro
195 200 205

Glu Asn Ile Leu Leu Asn Glu Asp Met His Ile Gln Ile Thr Asp Phe
210 215 220

Gly Thr Ala Lys Val Leu Ser Pro Glu Ser Lys Gln Val Cys
225 230 235

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<210> 155
<211> 73
<212> PRT
<213> Homo sapiens
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<400> 155

Met Ser Asp Val Thr Ile Val Lys Glu Gly Trp Val Gln Lys Arg Gly
1 5 10 15

Glu Tyr Ile Lys Asn Trp Arg Pro Arg Tyr Phe Leu Leu Lys Thr Asp
20 25 30

Gly Ser Phe Ile Gly Tyr Lys Glu Lys Pro Gln Asp Val Asp Leu Pro
35 40 45

Tyr Pro Leu Asn Asn Phe Ser Val Ala Ser Ser Val Met Phe Arg Tyr
50 55 60

Leu Gln Asn Leu Thr Leu Asn Gln Val
65 70

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<210> 156
<211> 213
<212> PRT
<213> Homo sapiens
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<400> 156

Met Ser Asp Val Thr Ile Val Lys Glu Gly Trp Val Gln Lys Arg Gly
1 5 10 15

Glu Tyr Ile Lys Asn Trp Arg Pro Arg Tyr Phe Leu Leu Lys Thr Asp
20 25 30

Gly Ser Phe Ile Gly Tyr Lys Glu Lys Pro Gln Asp Val Asp Leu Pro
35 40 45

Tyr Pro Leu Asn Asn Phe Ser Val Ala Lys Cys Gln Leu Met Lys Thr
50 55 60

Glu Arg Pro Lys Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp Thr
65 70 75 80

Thr Val Ile Glu Arg Thr Phe His Val Asp Thr Pro Glu Glu Arg Glu
85 90 95

Glu Trp Thr Glu Ala Ile Gln Ala Val Ala Asp Arg Leu Gln Arg Gln
100 105 110

Glu Glu Glu Arg Met Asn Cys Ser Pro Thr Ser Gln Ile Asp Asn Ile
115 120 125

Gly Glu Glu Glu Met Asp Ala Ser Thr Thr His His Lys Arg Lys Thr
130 135 140

Met Asn Asp Phe Asp Tyr Leu Lys Leu Leu Gly Lys Gly Thr Phe Gly
145 150 155 160

Lys Val Ile Leu Val Arg Glu Lys Ala Ser Gly Lys Tyr Tyr Ala Met
165 170 175

Lys Ile Leu Lys Lys Glu Val Ile Ile Ala Lys Val Thr Asp Leu Leu
180 185 190

Lys Leu Ile Thr Lys Phe Leu Phe Ala Val Cys Met Cys Leu Trp Ala
195 200 205

His Glu Phe Thr Cys
210

<210> 157

<211> 352

<212> PRT

<213> Homo sapiens

<400> 157

Met Gly Gly Lys Pro Ala Asn Arg Met Met Pro Tyr Pro Phe Pro Ser
1 5 10 15

Gly Thr Trp Lys Val Lys Trp Val Ala Ser Arg Asn Ala Phe Lys Pro
20 25 30

Arg	Ile	Gly	Ile	Leu	Ile	Lys	Thr	Leu	Ile	Tyr	Ser	Ser	Gln	Phe	Pro
		35					40					45			
Leu	Gly	Asn	Leu	Glu	Lys	Ile	Ser	Gln	Leu	Leu	Ser	Lys	Ser	Ala	Gln
	50					55					60				
Cys	Pro	Leu	Arg	Val	His	Tyr	Leu	Ser	Ser	Gln	Tyr	Gly	Asp	Glu	Arg
65					70					75					80
Cys	Phe	Met	Phe	Val	Leu	Ile	Ser	Pro	Thr	Lys	Ser	Val	Ile	Ile	Thr
				85					90					95	
Ile	Leu	Ser	Leu	Leu	Phe	Thr	Leu	Gln	Leu	Phe	Phe	His	Leu	Ser	Arg
			100					105					110		
Glu	Arg	Val	Phe	Ser	Glu	Asp	Arg	Thr	Arg	Phe	Tyr	Gly	Ala	Glu	Ile
		115					120					125			
Val	Ser	Ala	Leu	Asp	Tyr	Leu	His	Ser	Gly	Lys	Ile	Val	Tyr	Arg	Asp
		130				135					140				
Leu	Lys	Leu	Glu	Asn	Leu	Met	Leu	Asp	Lys	Asp	Gly	His	Ile	Lys	Ile
145				150						155					160
Thr	Asp	Phe	Gly	Leu	Cys	Lys	Glu	Gly	Ile	Thr	Asp	Ala	Ala	Thr	Met
				165					170					175	
Lys	Thr	Phe	Cys	Gly	Thr	Pro	Glu	Tyr	Leu	Ala	Pro	Glu	Val	Leu	Glu
			180					185					190		
Asp	Asn	Asp	Tyr	Gly	Arg	Ala	Val	Asp	Trp	Trp	Gly	Leu	Gly	Val	Val
		195					200					205			
Met	Tyr	Glu	Met	Met	Cys	Gly	Arg	Leu	Pro	Phe	Tyr	Asn	Gln	Asp	His
		210				215					220				
Glu	Lys	Leu	Phe	Glu	Leu	Ile	Leu	Met	Glu	Asp	Ile	Lys	Phe	Pro	Arg
225					230					235					240
Thr	Leu	Ser	Ser	Asp	Ala	Lys	Ser	Leu	Leu	Ser	Gly	Leu	Leu	Ile	Lys
				245					250					255	
Asp	Pro	Asn	Lys	Arg	Leu	Gly	Gly	Gly	Pro	Asp	Asp	Ala	Lys	Glu	Ile
			260				265						270		
Met	Arg	His	Ser	Phe	Phe	Ser	Gly	Val	Asn	Trp	Gln	Asp	Val	Tyr	Asp
		275					280					285			
Lys	Lys	Leu	Val	Pro	Pro	Phe	Lys	Pro	Gln	Val	Thr	Ser	Glu	Thr	Asp
		290				295					300				
Thr	Arg	Tyr	Phe	Asp	Glu	Glu	Phe	Thr	Ala	Gln	Thr	Ile	Thr	Ile	Thr
305					310					315					320
Pro	Pro	Glu	Lys	Tyr	Asp	Glu	Asp	Gly	Met	Asp	Cys	Met	Asp	Asn	Glu
				325					330					335	

Arg Arg Pro His Phe Pro Gln Phe Ser Tyr Ser Ala Ser Gly Arg Glu
 340 345 350

<210> 158
 <211> 132
 <212> PRT
 <213> Homo sapiens

<400> 158

Met Glu Leu Leu Arg Thr Ile Thr Tyr Gln Pro Ala Ala Ser Thr Lys
 1 5 10 15

Met Cys Glu Gln Ala Leu Gly Lys Gly Cys Gly Gly Asn Ser Lys Lys
 20 25 30

Lys Arg Pro Pro Gln Pro Pro Glu Glu Ser Gln Pro Pro Gln Ser Gln
 35 40 45

Ala Gln Val Pro Pro Ala Ala Pro His His His His His His Ser His
 50 55 60

Ser Gly Pro Glu Ile Ser Arg Ile Ile Val Asp Pro Thr Thr Gly Lys
 65 70 75 80

Arg Tyr Cys Arg Gly Lys Val Leu Gly Lys Gly Gly Phe Ala Lys Cys
 85 90 95

Tyr Glu Met Thr Asp Leu Thr Asn Asn Lys Val Tyr Ala Ala Lys Ile
 100 105 110

Ile Pro His Ser Arg Val Ala Lys Pro His Gln Arg Glu Lys Val Cys
 115 120 125

Met Thr Leu Glu
 130

<210> 159
 <211> 192
 <212> PRT
 <213> Homo sapiens

<400> 159

Met Glu Leu Leu Arg Thr Ile Thr Tyr Gln Pro Ala Ala Ser Thr Lys
 1 5 10 15

Met Cys Glu Gln Ala Leu Gly Lys Gly Cys Gly Gly Asn Ser Lys Lys
 20 25 30

Lys Arg Pro Pro Gln Pro Pro Glu Glu Ser Gln Pro Pro Gln Ser Gln
 35 40 45

Ala Gln Val Pro Pro Ala Ala Pro His His His His His His Ser His
 50 55 60

Ser Gly Pro Glu Ile Ser Arg Ile Ile Val Asp Pro Thr Thr Gly Lys

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Phe Tyr His Tyr Phe Glu Asp Lys Glu Asn Ile Tyr Ile Leu Leu Glu
145 150 155 160

Tyr Cys Ser Arg Arg Leu Gln Gly Ser Gln Lys Asn Asp Leu Glu Tyr
165 170 175

Val Glu Glu Asp Gly His Val Val Val Arg Lys Gln Phe Pro Cys Gly
180 185 190

Leu Leu Asp Trp Val Glu Pro Glu Gln Ala Lys Ala Tyr Ser Ser
195 200 205

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<210> 161
<211> 337
<212> PRT
<213> Homo sapiens
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<400> 161

Met Ser Asp Lys Asp Leu Arg Thr Ala Ala Ala Gly Gly Gly His Leu
1 5 10 15

Val Ala Ile Leu Thr Val Phe Ile Pro Gln Lys Asp Leu Val Glu Glu
20 25 30

Glu Ala Glu Glu Ala Gly Val Ala Leu Arg Ser Thr Gln Ser Thr Leu
35 40 45

Gln Ala Gly Leu Ala Ala Asp Ala Trp Ala Ala Pro Ile Ala Met Gln
50 55 60

Ile Tyr Lys Lys His Leu Asp Pro Arg Pro Gly Pro Cys His Leu Ser
65 70 75 80

Trp Ala Trp Ala Trp Ala Ser Trp Pro Ala Ala Ala Cys Thr Ala Gly
85 90 95

Pro Lys Gly Arg Pro Pro Met Thr Gln Val Tyr Glu Arg Leu Glu Lys
100 105 110

Leu Gln Ala Val Val Ala Gly Val Pro Gly His Leu Glu Ala Ala Ser
115 120 125

Cys Ile Pro Phe Pro Gln Glu Asn Ser Tyr Val Ser Ser Thr Gly Arg
130 135 140

Ala Ser Ala Gln Ala Ala Glu Gln Leu Gln Arg Gly Pro Asn Gln Pro
145 150 155 160

Val Glu Ser Asp Glu Ser Leu Gly Gly Leu Ser Ala Ala Leu Arg Ser
165 170 175

Trp His Leu Thr Pro Ser Cys Pro Leu Asp Pro Ala Pro Leu Arg Glu
180 185 190

Ala Gly Cys Pro Gln Gly Asp Thr Ala Gly Glu Ser Ser Trp Gly Ser
195 200 205

[illegible]

Gly	Pro	Gly	Ser	Arg	Pro	Thr	Ala	Val	Glu	Gly	Leu	Ala	Leu	Gly	Ser
210						215					220				
Ser	Ala	Ser	Ser	Ser	Ser	Glu	Pro	Pro	Gln	Ile	Ile	Ile	Asn	Pro	Ala
225					230					235					240
Arg	Gln	Lys	Met	Val	Gln	Lys	Leu	Ala	Leu	Tyr	Glu	Asp	Gly	Ala	Leu
			245						250					255	
Asp	Ser	Leu	Gln	Leu	Leu	Ser	Ser	Ser	Ser	Leu	Pro	Gly	Leu	Gly	Leu
		260						265					270		
Glu	Gln	Asp	Arg	Gln	Gly	Pro	Lys	Lys	Val	Met	Asn	Phe	Arg	Ala	Asp
		275					280					285			
Val	Phe	Thr	Trp	Ala	Asp	Pro	Pro	Asn	Pro	Glu	Val	Lys	Val	Leu	Met
	290					295					300				
Val	Arg	Ser	Ser	His	Gly	Ala	Arg	Val	Leu	Ser	Thr	Leu	Pro	Ala	Val
305					310					315					320
Gly	Val	Gly	Ala	His	Ala	Arg	Trp	Gly	Glu	Lys	Glu	Val	Ala	Leu	Leu
				325					330					335	

Phe

<210> 162
 <211> 122
 <212> PRT
 <213> Homo sapiens

<400> 162

Met	Gly	His	Ala	Leu	Cys	Val	Cys	Ser	Arg	Gly	Thr	Val	Ile	Ile	Asp
1				5					10					15	
Asn	Lys	Arg	Tyr	Leu	Phe	Ile	Gln	Lys	Leu	Gly	Glu	Gly	Gly	Phe	Ser
			20					25					30		
Tyr	Val	Asp	Leu	Val	Glu	Gly	Leu	His	Asp	Gly	His	Phe	Tyr	Ala	Leu
		35					40					45			
Lys	Arg	Ile	Leu	Cys	His	Glu	Gln	Gln	Asp	Arg	Glu	Glu	Ala	Gln	Arg
	50					55					60				
Glu	Ala	Asp	Met	His	Arg	Leu	Phe	Asn	His	Pro	Asn	Ile	Leu	Arg	Leu
65					70					75					80
Val	Ala	Tyr	Cys	Leu	Arg	Glu	Arg	Gly	Ala	Lys	His	Glu	Ala	Trp	Leu
				85					90					95	
Leu	Leu	Pro	Phe	Phe	Lys	Val	Arg	Lys	Thr	Pro	Val	Tyr	Gly	Gly	Gly
			100					105					110		
Cys	Ser	Arg	Ala	Thr	Tyr	Ser	Arg	Ala	Val						

115

120

<210> 163
 <211> 842
 <212> PRT
 <213> Homo sapiens

<400> 163

Met	Glu	Arg	Ala	Ile	Ser	Pro	Gly	Leu	Leu	Val	Arg	Ala	Leu	Leu	Leu
1				5					10					15	
Leu	Leu	Leu	Leu	Gly	Leu	Ala	Ala	Arg	Thr	Val	Ala	Ala	Gly	Arg	Ala
			20					25					30		
Arg	Gly	Leu	Pro	Ala	Pro	Thr	Ala	Glu	Ala	Ala	Phe	Gly	Leu	Gly	Ala
		35					40					45			
Ala	Ala	Ala	Pro	Thr	Ser	Ala	Thr	Arg	Val	Pro	Ala	Ala	Gly	Ala	Val
		50				55					60				
Ala	Ala	Ala	Glu	Val	Thr	Val	Glu	Asp	Ala	Glu	Ala	Leu	Pro	Ala	Ala
65					70				75					80	
Ala	Gly	Glu	Gln	Glu	Pro	Arg	Gly	Pro	Glu	Pro	Asp	Asp	Glu	Thr	Glu
			85					90						95	
Leu	Arg	Pro	Arg	Gly	Arg	Ser	Leu	Val	Ile	Ile	Ser	Thr	Leu	Asp	Gly
		100						105					110		
Arg	Ile	Ala	Ala	Leu	Asp	Pro	Glu	Asn	His	Gly	Lys	Lys	Gln	Trp	Asp
		115					120					125			
Leu	Asp	Val	Gly	Ser	Gly	Ser	Leu	Val	Ser	Ser	Ser	Leu	Ser	Lys	Pro
	130					135					140				
Glu	Val	Phe	Gly	Asn	Lys	Met	Ile	Ile	Pro	Ser	Leu	Asp	Gly	Ala	Leu
145					150					155				160	
Phe	Gln	Trp	Asp	Arg	Asp	Arg	Glu	Ser	Met	Glu	Thr	Val	Pro	Phe	Thr
			165						170					175	
Val	Glu	Ser	Leu	Leu	Glu	Ser	Ser	Tyr	Lys	Phe	Gly	Asp	Asp	Val	Val
		180						185					190		
Leu	Val	Gly	Gly	Lys	Ser	Leu	Thr	Thr	Tyr	Gly	Leu	Ser	Ala	Tyr	Ser
		195					200					205			
Gly	Lys	Val	Arg	Tyr	Ile	Cys	Ser	Ala	Leu	Gly	Cys	Arg	Gln	Trp	Asp
	210					215					220				
Ser	Asp	Glu	Met	Glu	Gln	Glu	Glu	Asp	Ile	Leu	Leu	Leu	Gln	Arg	Thr
225					230					235				240	
Gln	Lys	Thr	Val	Arg	Ala	Val	Gly	Pro	Arg	Ser	Gly	Asn	Glu	Lys	Trp
			245						250					255	

Asn	Phe	Ser	Val	Gly	His	Phe	Glu	Leu	Arg	Tyr	Ile	Pro	Asp	Met	Glu	260	265	270
Thr	Arg	Ala	Gly	Phe	Ile	Glu	Ser	Thr	Phe	Lys	Pro	Asn	Glu	Asn	Thr	275	280	285
Glu	Glu	Ser	Lys	Ile	Ile	Ser	Asp	Val	Glu	Glu	Gln	Glu	Ala	Ala	Ile	290	295	300
Met	Asp	Ile	Val	Ile	Lys	Val	Ser	Val	Ala	Asp	Trp	Lys	Val	Met	Ala	305	310	315
Phe	Ser	Lys	Lys	Gly	Gly	His	Leu	Glu	Trp	Glu	Tyr	Gln	Phe	Cys	Thr	325	330	335
Pro	Ile	Ala	Ser	Ala	Trp	Leu	Leu	Lys	Asp	Gly	Lys	Val	Ile	Pro	Ile	340	345	350
Ser	Leu	Phe	Asp	Asp	Thr	Ser	Tyr	Thr	Ser	Asn	Asp	Asp	Val	Leu	Glu	355	360	365
Asp	Glu	Glu	Asp	Ile	Val	Glu	Ala	Ala	Arg	Gly	Ala	Thr	Glu	Asn	Ser	370	375	380
Val	Tyr	Leu	Gly	Met	Tyr	Arg	Gly	Gln	Leu	Tyr	Leu	Gln	Ser	Ser	Val	385	390	395
Arg	Ile	Ser	Glu	Lys	Phe	Pro	Ser	Ser	Pro	Lys	Ala	Leu	Glu	Ser	Val	405	410	415
Thr	Asn	Glu	Asn	Ala	Ile	Ile	Pro	Leu	Pro	Thr	Ile	Lys	Trp	Lys	Pro	420	425	430
Leu	Ile	His	Ser	Pro	Ser	Arg	Thr	Pro	Val	Leu	Val	Gly	Ser	Asp	Glu	435	440	445
Phe	Asp	Lys	Cys	Leu	Ser	Asn	Asp	Lys	Phe	Ser	His	Glu	Glu	Tyr	Ser	450	455	460
Asn	Gly	Ala	Leu	Ser	Ile	Leu	Gln	Tyr	Pro	Tyr	Asp	Asn	Gly	Tyr	Tyr	465	470	475
Leu	Pro	Tyr	Tyr	Lys	Arg	Glu	Arg	Asn	Lys	Arg	Ser	Thr	Gln	Ile	Thr	485	490	495
Val	Arg	Phe	Leu	Asp	Asn	Pro	His	Tyr	Asn	Lys	Asn	Ile	Arg	Lys	Lys	500	505	510
Asp	Pro	Val	Leu	Leu	Leu	His	Trp	Trp	Lys	Glu	Ile	Val	Ala	Thr	Ile	515	520	525
Leu	Phe	Cys	Ile	Ile	Ala	Thr	Thr	Phe	Ile	Val	Arg	Arg	Leu	Phe	His	530	535	540
Pro	His	Pro	His	Arg	Gln	Arg	Lys	Glu	Ser	Glu	Thr	Gln	Cys	Gln	Thr	545	550	555

Glu Asn Lys Tyr Asp Ser Val Ser Gly Glu Ala Asn Asp Ser Ser Trp
 565 570 575
 Asn Asp Ile Lys Asn Ser Gly Tyr Ile Ser Arg Tyr Leu Thr Asp Phe
 580 585 590
 Glu Pro Ile Gln Cys Leu Gly Arg Gly Gly Phe Gly Val Val Phe Glu
 595 600 605
 Ala Lys Asn Lys Val Asp Asp Cys Asn Tyr Ala Ile Lys Arg Ile Arg
 610 615 620
 Leu Pro Asn Arg Glu Leu Ala Arg Glu Lys Val Met Arg Glu Val Lys
 625 630 635 640
 Ala Leu Ala Lys Leu Glu His Pro Gly Ile Val Arg Tyr Phe Asn Ala
 645 650 655
 Trp Leu Glu Ala Pro Pro Glu Lys Trp Gln Glu Lys Met Asp Glu Ile
 660 665 670
 Trp Leu Lys Asp Glu Ser Thr Asp Trp Pro Leu Ser Ser Pro Ser Pro
 675 680 685
 Met Asp Ala Pro Ser Val Lys Ile Arg Arg Met Asp Pro Phe Ser Thr
 690 695 700
 Lys Glu His Ile Glu Ile Ile Ala Pro Ser Pro Gln Arg Ser Arg Ser
 705 710 715 720
 Phe Ser Val Gly Ile Ser Cys Asp Gln Thr Ser Ser Ser Glu Ser Gln
 725 730 735
 Phe Ser Pro Leu Glu Phe Ser Gly Met Asp His Glu Asp Ile Ser Glu
 740 745 750
 Ser Val Asp Ala Ala Tyr Asn Leu Gln Asp Ser Cys Leu Thr Asp Cys
 755 760 765
 Asp Val Glu Asp Gly Thr Met Asp Gly Asn Asp Glu Gly His Ser Phe
 770 775 780
 Glu Leu Cys Pro Ser Glu Ala Ser Pro Tyr Val Arg Ser Arg Glu Arg
 785 790 795 800
 Thr Ser Ser Ser Ile Val Phe Glu Asp Ser Gly Cys Asp Asn Ala Ser
 805 810 815
 Ser Lys Glu Glu Pro Lys Thr Asn Arg Leu His Ile Gly Asn His Cys
 820 825 830
 Ala Asn Lys Leu Thr Val Thr Val Leu Phe
 835 840

<210> 164

<211> 743

<212> PRT

[illegible]

Met 1	Gly	Ser	Arg	Ala 5	Gln	Lys	Ser	Ala	Gly 10	Asn	Ala	Glu	Leu	Trp 15	Glu
Pro	Leu	Pro	Glu 20	Gly	Arg	Pro	Arg	Pro	Ala 25	Gly	Thr	Ser	Ser	Ala	Val
Ser	Ala	Trp 35	Ala	Ser	Leu	Lys	Leu	Cys	Leu	Arg	Gly	Gly 45	Ser	Gly	Arg
Arg	Gln 50	Arg	Leu	Gly	Gly	Gly 55	Arg	Met	Gln	Pro	Glu 60	Glu	Gly	His	Arg
Leu 65	Ala	Ala	Gly	Ala	Ala 70	Val	Arg	Gly	Ala	Ala 75	Ala	Thr	Val	Leu	Leu 80
Arg	Leu	Arg	Asp 85	Asp	Leu	Asn	Val	Thr	Arg 90	Leu	Ser	His	Phe	Glu 95	Tyr
Val	Lys	Asn 100	Glu	Asp	Leu	Glu	Lys	Ile	Gly 105	Met	Gly	Arg	Pro 110	Gly	Gln
Arg	Arg	Leu 115	Trp	Glu	Ala	Val	Lys	Arg	Arg	Lys	Ala	Leu 125	Cys	Lys	Arg
Lys 130	Ser	Trp	Met	Asn	Lys	Val 135	Phe	Ser	Gly	Lys	Arg 140	Leu	Glu	Ala	Glu
Phe 145	Pro	Pro	His	His	Ser 150	Gln	Ser	Thr	Phe	Arg 155	Lys	Thr	Ser	Pro	Ala 160
Pro	Gly	Gly	Pro	Ala 165	Gly	Glu	Gly	Pro	Leu 170	Gln	Ser	Leu	Thr	Cys 175	Leu
Ile	Gly	Glu 180	Lys	Asp	Leu	Arg	Leu	Leu 185	Glu	Lys	Leu	Gly 190	Asp	Gly	Ser
Phe 195	Gly	Val	Val	Arg	Arg	Gly	Glu 200	Trp	Asp	Ala	Pro 205	Ser	Gly	Lys	Thr
Val 210	Ser	Pro	Pro	Gln	Pro	Ala 215	Phe	Phe	Thr	Gln	Lys 220	Pro	Thr	Tyr	Asp
Pro 225	Val	Ser	Glu	Asp	Gln 230	Asp	Pro	Leu	Ser	Ser 235	Asp	Phe	Lys	Arg	Leu 240
Gly	Leu	Arg	Lys	Pro 245	Gly	Leu	Pro	Arg	Gly 250	Leu	Trp	Leu	Ala	Lys 255	Pro
Ser	Ala	Arg	Val 260	Pro	Gly	Thr	Lys	Ala 265	Ser	Arg	Gly	Ser	Gly 270	Ala	Glu
Val 275	Thr	Leu	Ile	Asp	Phe	Gly	Glu 280	Glu	Pro	Val	Val 285	Pro	Ala	Leu	Arg

Pro Cys Ala Pro Ser Leu Ala Gln Leu Ala Met Asp Ala Cys Ser Leu
290 295 300

Leu Asp Glu Thr Pro Pro Gln Ser Pro Thr Arg Ala Leu Pro Arg Pro
305 310 315 320

Leu His Pro Thr Pro Val Val Asp Trp Asp Ala Arg Pro Leu Pro Pro
325 330 335

Pro Pro Ala Tyr Asp Asp Val Ala Gln Asp Glu Asp Asp Phe Glu Ile
340 345 350

Cys Ser Ile Asn Ser Thr Leu Val Gly Ala Gly Val Pro Ala Gly Pro
355 360 365

Ser Gln Gly Gln Thr Asn Tyr Ala Phe Val Pro Glu Gln Ala Arg Pro
370 375 380

Pro Pro Pro Leu Glu Asp Asn Leu Phe Leu Pro Pro Gln Gly Gly Gly
385 390 395 400

Lys Pro Pro Ser Ser Ala Gln Thr Ala Glu Ile Phe Gln Ala Leu Gln
405 410 415

Gln Glu Cys Met Arg Gln Leu Gln Ala Pro Ala Gly Ser Pro Ala Pro
 420 425 430

Ser	Pro	Ser	Pro	Gly	Gly	Asp	Asp	Lys	Pro	Gln	Val	Pro	Pro	Arg	Val
		435					440					445			

Pro Ile Pro Pro Arg Pro Thr Arg Pro His Val Gln Leu Ser Pro Ala
450 455 460

Pro Pro Gly Glu Glu Glu Thr Ser Gln Trp Pro Gly Pro Ala Ser Pro
465 470 475 480

Pro Arg Val Pro Pro Arg Glu Pro Leu Ser Pro Gln Gly Ser Arg Thr
485 490 495

Pro Ser Pro Leu Val Pro Pro Gly Ser Ser Pro Leu Pro Pro Arg Leu
500 505 510

Ser Ser Ser Pro Gly Lys Thr Met Pro Thr Thr Gln Ser Phe Ala Ser
515 520 525

Asp Pro Lys Tyr Ala Thr Pro Gln Val Ile Gln Ala Pro Gly Pro Arg
530 535 540

Ala Gly Pro Cys Ile Leu Pro Ile Val Arg Asp Gly Lys Lys Val Ser
545 550 555 560

Ser Thr His Tyr Tyr Leu Leu Pro Glu Arg Pro Ser Tyr Leu Glu Arg
565 570 575

Tyr Gln Arg Phe Leu Arg Glu Ala Gln Ser Pro Glu Glu Pro Thr Pro
580 585 590

Leu Pro Val Pro Leu Leu Leu Pro Pro Pro Ser Thr Pro Ala Pro Ala
595 600 605

Ala Pro Thr Ala Thr Val Arg Pro Met Pro Gln Ala Ala Leu Asp Pro
610 615 620

Lys Ala Asn Phe Ser Thr Asn Asn Ser Asn Pro Gly Ala Arg Pro Pro
625 630 635 640

Pro Pro Arg Ala Thr Ala Arg Leu Pro Gln Arg Gly Cys Pro Gly Asp
645 650 655

Gly Pro Glu Ala Gly Arg Pro Ala Asp Lys Ile Gln Met Ala Met Val
660 665 670

His Gly Val Thr Thr Glu Glu Cys Gln Ala Ala Leu Gln Cys His Gly
675 680 685

Trp Ser Val Gln Arg Ala Cys Pro Val Ser Glu Gly Gly Ala Ala Leu
690 695 700

Arg Ala Gly Ser Ala Ala Gln Arg Glu Cys His Lys Val Leu Glu Met
705 710 715 720

Phe Asp Trp Asn Leu Glu Gln Ala Gly Cys His Leu Leu Gly Ser Trp
725 730 735

Gly Pro Ala His His Lys Arg
740

<210> 165
<211> 604
<212> PRT
<213> Homo sapiens

<400> 165

Met Ala Ser Asn Pro Glu Arg Gly Glu Ile Leu Leu Thr Glu Leu Gln
1 5 10 15

Gly Asp Ser Arg Ser Leu Pro Phe Ser Glu Asn Val Ser Ala Val Gln
20 25 30

Lys Leu Asp Phe Ser Asp Thr Met Val Gln Gln Lys Leu Asp Asp Ile
35 40 45

Lys Asp Arg Ile Lys Arg Glu Ile Arg Lys Glu Leu Lys Ile Lys Glu
50 55 60

Gly Ala Glu Asn Leu Arg Lys Val Thr Thr Asp Lys Lys Ser Leu Ala
65 70 75 80

Tyr Val Asp Asn Ile Leu Lys Lys Ser Asn Lys Lys Leu Glu Glu Leu
85 90 95

His His Lys Leu Gln Glu Leu Asn Ala His Ile Val Val Ser Asp Pro

165
604
PRT
Homo sapiens
165

100					105					110						
Glu	Asp	Ile	Thr	Asp	Cys	Pro	Arg	Thr	Pro	Asp	Thr	Pro	Asn	Asn	Asp	
115					120					125						
Pro	Arg	Cys	Ser	Thr	Ser	Asn	Asn	Arg	Leu	Lys	Ala	Leu	Gln	Lys	Gln	
130					135					140						
Leu	Asp	Ile	Glu	Leu	Lys	Val	Lys	Gln	Gly	Ala	Glu	Asn	Met	Ile	Gln	
145					150					155					160	
Met	Tyr	Ser	Asn	Gly	Ser	Ser	Lys	Asp	Arg	Lys	Leu	His	Gly	Thr	Ala	
165					170					175						
Gln	Gln	Leu	Leu	Gln	Asp	Ser	Lys	Thr	Lys	Ile	Glu	Val	Ile	Arg	Met	
180					185					190						
Gln	Ile	Leu	Gln	Ala	Val	Gln	Thr	Asn	Glu	Leu	Ala	Phe	Asp	Asn	Ala	
195					200					205						
Lys	Pro	Val	Ile	Ser	Pro	Leu	Glu	Leu	Arg	Met	Glu	Glu	Leu	Arg	His	
210					215					220						
His	Phe	Arg	Ile	Glu	Phe	Ala	Val	Ala	Glu	Gly	Ala	Lys	Asn	Val	Met	
225					230					235					240	
Lys	Leu	Leu	Gly	Ser	Gly	Lys	Val	Thr	Asp	Arg	Lys	Ala	Leu	Ser	Glu	
245					250					255						
Ala	Gln	Ala	Arg	Phe	Asn	Glu	Ser	Ser	Gln	Lys	Leu	Asp	Leu	Leu	Lys	
260					265					270						
Tyr	Ser	Leu	Glu	Gln	Arg	Leu	Asn	Glu	Val	Pro	Lys	Asn	His	Pro	Lys	
275					280					285						
Ser	Arg	Ile	Ile	Ile	Glu	Glu	Leu	Ser	Leu	Val	Ala	Ala	Ser	Pro	Thr	
290					295					300						
Leu	Ser	Pro	Arg	Gln	Ser	Met	Ile	Ser	Thr	Gln	Asn	Gln	Tyr	Ser	Thr	
305					310					315					320	
Leu	Ser	Lys	Pro	Ala	Ala	Leu	Thr	Gly	Thr	Leu	Glu	Val	Arg	Leu	Met	
325					330					335						
Gly	Cys	Gln	Asp	Ile	Leu	Glu	Asn	Val	Pro	Gly	Arg	Ser	Lys	Ala	Thr	
340					345					350						
Ser	Val	Ala	Leu	Pro	Gly	Trp	Ser	Pro	Ser	Glu	Thr	Arg	Ser	Ser	Phe	
355					360					365						
Met	Ser	Arg	Thr	Ser	Lys	Ser	Lys	Ser	Gly	Ser	Ser	Arg	Asn	Leu	Leu	
370					375					380						
Lys	Thr	Asp	Asp	Leu	Ser	Asn	Asp	Val	Cys	Ala	Val	Leu	Lys	Leu	Asp	
385					390					395					400	
Asn	Thr	Val	Val	Gly	Gln	Thr	Ser	Trp	Lys	Pro	Ile	Ser	Asn	Gln	Ser	

100 105 110
 115 120 125
 130 135 140
 145 150 155 160
 165 170 175
 180 185 190
 195 200 205
 210 215 220
 225 230 235 240
 245 250 255
 260 265 270
 275 280 285
 290 295 300
 305 310 315 320
 325 330 335
 340 345 350
 355 360 365
 370 375 380
 385 390 395 400

405										410					415				
Trp	Asp	Gln	Lys	Phe	Thr	Leu	Glu	Leu	Asp	Arg	Ser	Arg	Glu	Leu	Glu				
			420						425				430						
Ile	Ser	Val	Tyr	Trp	Arg	Asp	Trp	Arg	Ser	Leu	Cys	Ala	Val	Lys	Phe				
		435					440					445							
Leu	Arg	Leu	Glu	Asp	Phe	Leu	Asp	Asn	Gln	Arg	His	Gly	Met	Cys	Leu				
		450				455					460								
Tyr	Leu	Glu	Pro	Gln	Gly	Thr	Leu	Phe	Ala	Glu	Val	Thr	Phe	Phe	Asn				
465					470					475					480				
Pro	Val	Ile	Glu	Arg	Arg	Pro	Lys	Leu	Gln	Arg	Gln	Lys	Lys	Ile	Phe				
				485				490						495					
Ser	Lys	Gln	Gln	Gly	Lys	Thr	Phe	Leu	Arg	Ala	Pro	Gln	Met	Asn	Ile				
			500					505					510						
Asn	Ile	Ala	Thr	Trp	Gly	Arg	Leu	Val	Arg	Arg	Ala	Ile	Pro	Thr	Val				
		515					520					525							
Asn	His	Ser	Gly	Thr	Phe	Ser	Pro	Gln	Ala	Pro	Val	Pro	Thr	Thr	Val				
	530					535					540								
Pro	Val	Val	Asp	Val	Arg	Ile	Pro	Gln	Leu	Ala	Pro	Pro	Ala	Arg	Tyr				
545					550					555					560				
Val	Ser	Glu	Ile	Leu	Ser	Ile	Ser	Tyr	Thr	Lys	Leu	Leu	Gly	His	Ser				
			565					570						575					
Tyr	Val	Leu	Ile	Ile	Ala	Gly	Val	Leu	Ser	Leu	Ala	Phe	Phe	Pro	Ser				
		580						585					590						
Ser	Ile	Leu	Lys	Val	Val	Phe	Cys	Leu	Leu	Lys	Lys								
		595					600												

<210> 166
 <211> 613
 <212> PRT
 <213> Homo sapiens

<400> 166

Met	Ala	Ser	Asn	Pro	Glu	Arg	Gly	Glu	Ile	Leu	Leu	Thr	Glu	Leu	Gln
1			5					10					15		
Gly	Asp	Ser	Arg	Ser	Leu	Pro	Phe	Ser	Glu	Asn	Val	Ser	Ala	Val	Gln
		20					25					30			
Lys	Leu	Asp	Phe	Ser	Asp	Thr	Met	Val	Gln	Gln	Lys	Leu	Asp	Asp	Ile
		35				40						45			
Lys	Asp	Arg	Ile	Lys	Arg	Glu	Ile	Arg	Lys	Glu	Leu	Lys	Ile	Lys	Glu
	50				55					60					

Gly	Ala	Glu	Asn	Leu	Arg	Lys	Val	Thr	Thr	Asp	Lys	Lys	Ser	Leu	Ala	
65					70					75					80	
Tyr	Val	Asp	Asn	Ile	Leu	Lys	Lys	Ser	Asn	Lys	Lys	Leu	Glu	Glu	Leu	
			85						90					95		
His	His	Lys	Leu	Gln	Glu	Leu	Asn	Ala	His	Ile	Val	Val	Ser	Asp	Pro	
			100					105						110		
Glu	Asp	Ile	Thr	Asp	Cys	Pro	Arg	Thr	Pro	Asp	Thr	Pro	Asn	Asn	Asp	
		115					120						125			
Pro	Arg	Cys	Ser	Thr	Ser	Asn	Asn	Arg	Leu	Lys	Ala	Leu	Gln	Lys	Gln	
	130					135					140					
Leu	Asp	Ile	Glu	Leu	Lys	Val	Lys	Gln	Gly	Ala	Glu	Asn	Met	Ile	Gln	
145					150					155					160	
Met	Tyr	Ser	Asn	Gly	Ser	Ser	Lys	Asp	Arg	Lys	Leu	His	Gly	Thr	Ala	
			165						170					175		
Gln	Gln	Leu	Leu	Gln	Asp	Ser	Lys	Thr	Lys	Ile	Glu	Val	Ile	Arg	Met	
		180						185					190			
Gln	Ile	Leu	Gln	Ala	Val	Gln	Thr	Asn	Glu	Leu	Ala	Phe	Asp	Asn	Ala	
	195						200					205				
Lys	Pro	Val	Ile	Ser	Pro	Leu	Glu	Leu	Arg	Met	Glu	Glu	Leu	Arg	His	
	210					215					220					
His	Phe	Arg	Ile	Glu	Phe	Ala	Val	Ala	Glu	Gly	Ala	Lys	Asn	Val	Met	
225					230					235					240	
Lys	Leu	Leu	Gly	Ser	Gly	Lys	Val	Thr	Asp	Arg	Lys	Ala	Leu	Ser	Glu	
			245						250					255		
Ala	Gln	Ala	Arg	Phe	Asn	Glu	Ser	Ser	Gln	Lys	Leu	Asp	Leu	Leu	Lys	
			260					265					270			
Tyr	Ser	Leu	Glu	Gln	Arg	Leu	Asn	Glu	Val	Pro	Lys	Asn	His	Pro	Lys	
		275					280					285				
Ser	Arg	Ile	Ile	Ile	Glu	Glu	Leu	Ser	Leu	Val	Ala	Ala	Ser	Pro	Thr	
	290					295					300					
Leu	Ser	Pro	Arg	Gln	Ser	Met	Ile	Ser	Thr	Gln	Asn	Gln	Tyr	Ser	Thr	
305					310					315					320	
Leu	Ser	Lys	Pro	Ala	Ala	Leu	Thr	Gly	Thr	Leu	Glu	Val	Arg	Leu	Met	
			325						330					335		
Gly	Cys	Gln	Asp	Ile	Leu	Glu	Asn	Val	Pro	Gly	Arg	Ser	Lys	Ala	Thr	
			340					345					350			
Ser	Val	Ala	Leu	Pro	Gly	Trp	Ser	Pro	Ser	Glu	Thr	Arg	Ser	Ser	Phe	
		355					360					365				

Met Ser Arg Thr Ser Lys Ser Lys Ser Gly Ser Ser Arg Asn Leu Leu
370 375 380

Lys Thr Asp Asp Leu Ser Asn Asp Val Cys Ala Val Leu Lys Leu Asp
385 390 395 400

Asn Thr Val Val Gly Gln Thr Ser Trp Lys Pro Ile Ser Asn Gln Ser
405 410 415

Trp Asp Gln Lys Phe Thr Leu Glu Leu Asp Arg Ser Arg Glu Leu Glu
420 425 430

Ile Ser Val Tyr Trp Arg Asp Trp Arg Ser Leu Cys Ala Val Lys Phe
435 440 445

Leu Arg Leu Glu Asp Phe Leu Asp Asn Gln Arg His Gly Met Cys Leu
450 455 460

Tyr Leu Glu Pro Gln Gly Thr Leu Phe Ala Glu Val Thr Phe Phe Asn
465 470 475 480

Pro Val Ile Glu Arg Arg Pro Lys Leu Gln Arg Gln Lys Lys Ile Phe
485 490 495

Ser Lys Gln Gln Gly Lys Thr Phe Leu Arg Ala Pro Gln Met Asn Ile
500 505 510

Asn Ile Ala Thr Trp Gly Arg Leu Val Arg Arg Ala Ile Pro Thr Val
515 520 525

Asn His Ser Gly Thr Phe Ser Pro Gln Ala Pro Val Pro Thr Thr Val
530 535 540

Pro Val Val Asp Val Arg Ile Pro Gln Leu Ala Pro Pro Ala Ser Asp
545 550 555 560

Ser Thr Val Thr Lys Leu Asp Phe Asp Leu Glu Pro Glu Pro Pro Pro
565 570 575

Ala Pro Pro Arg Ala Ser Ser Leu Gly Glu Ile Asp Glu Ser Ser Glu
580 585 590

Leu Arg Val Leu Asp Ile Pro Gly Gln Ala Ser His Phe Lys Pro Cys
595 600 605

Ile Ile Pro Leu His
610

<210> 167

<211> 133

<212> PRT

<213> Homo sapiens

<400> 167

Met Val Ser Ser Gln Lys Leu Glu Lys Pro Ile Glu Met Gly Ser Ser
1 5 10 15

Glu Pro Leu Pro Ile Ala Asp Gly Asp Arg Arg Arg Lys Lys Lys Arg
20 25 30

Arg Gly Arg Ala Thr Asp Ser Leu Pro Gly Lys Phe Glu Asp Met Tyr
35 40 45

Lys Leu Thr Ser Glu Leu Leu Gly Glu Gly Ala Tyr Ala Lys Val Gln
50 55 60

Gly Ala Val Ser Leu Gln Asn Gly Lys Glu Tyr Ala Val Lys Ile Ile
65 70 75 80

Glu Lys Gln Ala Gly His Ser Arg Ser Arg Val Phe Arg Glu Val Glu
85 90 95

Thr Leu Tyr Gln Cys Gln Gly Asn Lys Asn Ile Leu Glu Leu Ile Glu
100 105 110

Phe Phe Glu Asp Asp Thr Arg Phe Tyr Leu Val Phe Glu Lys Leu Gln
115 120 125

Gly Gly Thr Tyr Arg
130

<210> 168

<211> 153

<212> PRT

<213> Homo sapiens

<400> 168

Met Leu Gln Val Gly Val Leu Arg Asp Arg Ser Pro Ala Gly Ala Ser
1 5 10 15

Glu Gly Phe His Val Arg Gly Arg Trp Arg Thr Glu Asp Cys His Leu
20 25 30

Arg Thr Lys Ala Ile Glu Thr Leu Arg Val Ala Gly Arg His Gln Leu
35 40 45

Pro Asp Arg Ser Phe Ile Ser Phe Gly Ile Ser Ser Leu Gln Met Val
50 55 60

Ser Ser Gln Lys Leu Glu Lys Pro Ile Glu Met Gly Ser Ser Glu Pro
65 70 75 80

Leu Pro Ile Ala Asp Gly Asp Arg Arg Arg Lys Lys Lys Arg Arg Gly
85 90 95

Arg Ala Thr Asp Ser Leu Pro Gly Lys Phe Glu Asp Met Tyr Lys Leu
100 105 110

Thr Ser Glu Leu Leu Gly Glu Gly Ala Tyr Ala Lys Val Gln Gly Ala
115 120 125

Val Ser Leu Gln Asn Gly Lys Glu Tyr Ala Val Lys Val Ser Val Ser

1000
900
800
700
600
500
400
300
200
100
0

130	135	140
Ala Glu Cys Gln Ala Leu Leu Cys Lys		
145	150	
<210> 169		
<211> 231		
<212> PRT		
<213> Homo sapiens		
<400> 169		
Met Gly Ser Gly Met Lys Leu Asn Asn Ser Cys Thr Pro Ile Thr Thr		
1	5	10 15
Pro Glu Leu Thr Thr Pro Cys Gly Ser Ala Glu Tyr Met Ala Pro Glu		
	20	25 30
Val Val Glu Val Phe Thr Asp Gln Ala Thr Phe Tyr Asp Lys Arg Cys		
	35	40 45
Asp Leu Trp Ser Leu Gly Val Val Leu Tyr Ile Met Leu Ser Gly Tyr		
	50	55 60
Pro Pro Phe Val Gly His Cys Gly Ala Asp Cys Gly Trp Asp Arg Gly		
65	70	75 80
Glu Val Cys Arg Val Cys Gln Asn Lys Leu Phe Glu Ser Ile Gln Glu		
	85	90 95
Gly Lys Tyr Glu Phe Pro Asp Lys Asp Trp Ala His Ile Ser Ser Glu		
	100	105 110
Ala Lys Asp Leu Ile Ser Lys Leu Leu Val Arg Asp Ala Lys Gln Arg		
	115	120 125
Leu Ser Ala Ala Gln Val Leu Gln His Pro Trp Val Gln Gly Gln Ala		
	130	135 140
Pro Glu Lys Gly Leu Pro Thr Pro Gln Val Leu Gln Arg Asn Ser Ser		
145	150	155 160
Thr Met Asp Leu Thr Leu Phe Ala Ala Glu Ala Ile Ala Leu Asn Arg		
	165	170 175
Gln Leu Ser Gln His Glu Glu Asn Glu Leu Ala Glu Glu Pro Glu Ala		
	180	185 190
Leu Ala Asp Gly Leu Cys Ser Met Lys Leu Ser Pro Pro Cys Lys Ser		
	195	200 205
Arg Leu Ala Arg Arg Arg Ala Leu Ala Gln Ala Gly Arg Gly Glu Asp		
	210	215 220
Arg Ser Pro Pro Thr Ala Leu		
225	230	

<210> 170
 <211> 146
 <212> PRT
 <213> Homo sapiens

<400> 170

Met	Arg	Lys	Gly	Val	Leu	Lys	Asp	Pro	Glu	Ile	Ala	Asp	Leu	Phe	Tyr
1				5					10					15	
Lys	Asp	Asp	Pro	Glu	Glu	Leu	Phe	Ile	Gly	Leu	His	Glu	Ile	Gly	His
			20					25					30		
Gly	Ser	Phe	Gly	Ala	Val	Tyr	Phe	Ala	Thr	Asn	Ala	His	Thr	Ser	Glu
		35					40					45			
Val	Val	Ala	Ile	Lys	Lys	Met	Ser	Tyr	Ser	Gly	Lys	Gln	Thr	His	Glu
	50					55					60				
Lys	Trp	Gln	Asp	Ile	Leu	Lys	Glu	Val	Lys	Phe	Leu	Arg	Gln	Leu	Lys
65					70					75					80
His	Pro	Asn	Thr	Ile	Glu	Tyr	Lys	Gly	Cys	Tyr	Leu	Lys	Glu	His	Thr
			85						90					95	
Ala	Trp	Leu	Val	Met	Glu	Tyr	Cys	Leu	Gly	Ser	Ala	Ser	Asp	Leu	Leu
			100					105					110		
Glu	Val	His	Lys	Lys	Pro	Leu	Gln	Glu	Val	Glu	Ile	Ala	Ala	Ile	Thr
		115					120					125			
His	Gly	Ala	Leu	His	Gly	Leu	Ala	Tyr	Leu	His	Ser	His	Ala	Leu	Ile
	130					135					140				
His	Arg														
145															

<210> 171
 <211> 123
 <212> PRT
 <213> Homo sapiens

<400> 171

Met	Met	Glu	Glu	Leu	His	Ser	Leu	Asp	Pro	Arg	Arg	Gln	Glu	Leu	Leu
1				5					10					15	
Glu	Ala	Arg	Phe	Thr	Gly	Val	Gly	Val	Ser	Lys	Gly	Pro	Leu	Asn	Ser
			20					25					30		
Glu	Ser	Ser	Asn	Gln	Ser	Leu	Cys	Ser	Val	Gly	Ser	Leu	Ser	Asp	Lys
		35					40					45			
Glu	Val	Glu	Thr	Pro	Glu	Lys	Lys	Gln	Asn	Asp	Gln	Arg	Asn	Arg	Lys
	50					55					60				
Arg	Lys	Ala	Glu	Pro	Tyr	Glu	Thr	Ser	Gln	Gly	Lys	Gly	Thr	Pro	Arg

65		70		75		80									
Gly	His	Lys	Ile	Ser	Asp	Tyr	Phe	Glu	Thr	Ala	Pro	Leu	Trp	Phe	Arg
				85					90					95	
Trp	Gln	Cys	Cys	Lys	Gly	Gly	Asn	Arg	Gly	Ala	Val	Cys	Ser	Ala	Asn
		100						105					110		
Pro	His	Val	Ser	Asp	Ala	Ser	Lys	Thr	Ser	Ala					
		115					120								

<210> 172
 <211> 478
 <212> PRT
 <213> Homo sapiens

<400> 172

Met	Val	Gly	Ile	Lys	Glu	Arg	Pro	Ser	Ser	Asn	Leu	Pro	Cys	Pro	Pro
1				5					10					15	
Leu	Pro	Pro	Gln	Thr	Gln	Ala	Cys	Pro	Pro	Leu	Ser	Trp	Pro	Gln	Arg
			20					25					30		
Leu	Asp	Ile	Leu	Leu	Gly	Thr	Ala	Arg	Ala	Ile	Gln	Phe	Leu	His	Gln
		35					40					45			
Asp	Ser	Pro	Ser	Leu	Ile	His	Gly	Asp	Ile	Lys	Ser	Ser	Asn	Val	Leu
	50					55					60				
Leu	Asp	Glu	Arg	Leu	Thr	Pro	Lys	Leu	Gly	Asp	Phe	Gly	Leu	Ala	Arg
65					70					75					80
Phe	Ser	Arg	Phe	Ala	Gly	Ser	Ser	Pro	Ser	Gln	Ser	Ser	Met	Val	Ala
				85					90					95	
Arg	Thr	Gln	Thr	Val	Arg	Gly	Thr	Leu	Ala	Tyr	Leu	Pro	Glu	Glu	Tyr
			100					105					110		
Ile	Lys	Thr	Gly	Arg	Leu	Ala	Val	Asp	Thr	Asp	Thr	Phe	Ser	Phe	Gly
		115					120					125			
Val	Val	Val	Leu	Glu	Thr	Leu	Ala	Gly	Gln	Arg	Ala	Val	Lys	Thr	His
		130					135					140			
Gly	Ala	Arg	Thr	Lys	Tyr	Leu	Lys	Asp	Leu	Val	Glu	Glu	Glu	Ala	Glu
145					150					155					160
Glu	Ala	Gly	Val	Ala	Leu	Arg	Ser	Thr	Gln	Ser	Thr	Leu	Gln	Ala	Gly
				165					170					175	
Leu	Ala	Ala	Asp	Ala	Trp	Ala	Ala	Pro	Ile	Ala	Met	Gln	Ile	Tyr	Lys
			180					185					190		
Lys	His	Leu	Asp	Pro	Arg	Pro	Gly	Pro	Cys	His	Leu	Ser	Trp	Ala	Trp
		195					200					205			

Ala	Trp	Ala	Ser	Trp	Pro	Ala	Ala	Ala	Cys	Thr	Ala	Gly	Pro	Lys	Gly	210	215	220	
Arg	Pro	Pro	Met	Thr	Gln	Val	Tyr	Glu	Arg	Leu	Glu	Lys	Leu	Gln	Ala	225	230	235	240
Val	Val	Ala	Gly	Val	Pro	Gly	His	Leu	Glu	Ala	Ala	Ser	Cys	Ile	Pro	245	250	255	
Phe	Pro	Gln	Glu	Asn	Ser	Tyr	Val	Ser	Ser	Thr	Gly	Arg	Ala	His	Ser	260	265	270	
Gly	Ala	Ala	Pro	Trp	Gln	Pro	Leu	Ala	Ala	Pro	Ser	Gly	Ala	Ser	Ala	275	280	285	
Gln	Ala	Ala	Glu	Gln	Leu	Gln	Arg	Gly	Pro	Asn	Gln	Pro	Val	Glu	Ser	290	295	300	
Asp	Glu	Ser	Leu	Gly	Gly	Leu	Ser	Ala	Ala	Leu	Arg	Ser	Trp	His	Leu	305	310	315	320
Thr	Pro	Ser	Cys	Pro	Leu	Asp	Pro	Ala	Pro	Leu	Arg	Glu	Ala	Gly	Cys	325	330	335	
Pro	Gln	Gly	Asp	Thr	Ala	Gly	Glu	Ser	Ser	Trp	Gly	Ser	Gly	Pro	Gly	340	345	350	
Ser	Arg	Pro	Thr	Ala	Val	Glu	Gly	Leu	Ala	Leu	Gly	Ser	Ser	Ala	Ser	355	360	365	
Ser	Ser	Ser	Glu	Pro	Pro	Gln	Ile	Ile	Ile	Asn	Pro	Ala	Arg	Gln	Lys	370	375	380	
Met	Val	Gln	Lys	Leu	Ala	Leu	Tyr	Glu	Asp	Gly	Ala	Leu	Asp	Ser	Leu	385	390	395	400
Gln	Leu	Leu	Ser	Ser	Ser	Ser	Leu	Pro	Gly	Leu	Gly	Leu	Glu	Gln	Asp	405	410	415	
Arg	Gln	Gly	Pro	Lys	Lys	Val	Met	Asn	Phe	Arg	Ala	Asp	Val	Phe	Thr	420	425	430	
Trp	Ala	Asp	Pro	Pro	Asn	Pro	Glu	Val	Lys	Val	Leu	Met	Val	Arg	Ser	435	440	445	
Ser	His	Gly	Ala	Arg	Val	Leu	Ser	Thr	Leu	Pro	Ala	Val	Gly	Val	Gly	450	455	460	
Ala	His	Ala	Arg	Trp	Gly	Glu	Lys	Glu	Val	Ala	Leu	Leu	Phe			465	470	475	

<210> 173

<211> 344

<212> PRT

<213> Homo sapiens

<400> 173

Met	Ala	Gly	Gly	Pro	Gly	Pro	Gly	Glu	Pro	Ala	Ala	Pro	Gly	Ala	Gln	1	5	10	15
His	Phe	Leu	Tyr	Glu	Val	Pro	Pro	Trp	Val	Met	Cys	Arg	Phe	Tyr	Lys	20	25	30	
Val	Met	Asp	Ala	Leu	Glu	Pro	Ala	Asp	Trp	Cys	Gln	Phe	Ala	Ala	Leu	35	40	45	
Ile	Val	Arg	Asp	Gln	Thr	Glu	Leu	Arg	Leu	Cys	Glu	Arg	Ser	Gly	Gln	50	55	60	
Arg	Thr	Ala	Ser	Val	Leu	Trp	Pro	Trp	Ile	Asn	Arg	Asn	Ala	Arg	Val	65	70	75	80
Ala	Asp	Leu	Val	His	Ile	Leu	Thr	His	Leu	Gln	Leu	Leu	Arg	Ala	Arg	85	90	95	
Asp	Ile	Ile	Thr	Ala	Trp	His	Pro	Pro	Ala	Pro	Leu	Pro	Ser	Pro	Gly	100	105	110	
Thr	Thr	Ala	Pro	Arg	Pro	Ser	Ser	Ile	Pro	Ala	Pro	Ala	Glu	Ala	Glu	115	120	125	
Ala	Trp	Ser	Pro	Arg	Lys	Leu	Pro	Ser	Ser	Ala	Ser	Thr	Phe	Leu	Ser	130	135	140	
Pro	Ala	Phe	Pro	Gly	Ser	Gln	Thr	His	Ser	Gly	Pro	Glu	Leu	Gly	Leu	145	150	155	160
Val	Pro	Ser	Pro	Ala	Ser	Leu	Trp	Pro	Pro	Pro	Pro	Ser	Pro	Ala	Pro	165	170	175	
Ser	Ser	Thr	Lys	Pro	Gly	Pro	Glu	Ser	Ser	Val	Ser	Leu	Leu	Gln	Gly	180	185	190	
Ala	Arg	Pro	Ser	Pro	Phe	Cys	Trp	Pro	Leu	Cys	Glu	Ile	Ser	Arg	Gly	195	200	205	
Thr	His	Asn	Phe	Ser	Glu	Glu	Leu	Lys	Ile	Gly	Glu	Gly	Gly	Phe	Gly	210	215	220	
Cys	Val	Tyr	Arg	Ala	Val	Met	Arg	Asn	Thr	Val	Tyr	Ala	Val	Lys	Arg	225	230	235	240
Leu	Lys	Glu	Asn	Ala	Asp	Leu	Glu	Trp	Thr	Ala	Val	Lys	Gln	Ser	Phe	245	250	255	
Leu	Thr	Glu	Val	Glu	Gln	Leu	Ser	Arg	Phe	Arg	His	Pro	Asn	Ile	Val	260	265	270	
Asp	Phe	Ala	Gly	Tyr	Cys	Ala	Gln	Asn	Gly	Phe	Tyr	Cys	Leu	Val	Tyr	275	280	285	
Gly	Phe	Leu	Pro	Asn	Gly	Ser	Leu	Glu	Asp	Arg	Leu	His	Cys	Gln	Thr	290	295	300	

210

215

220

Val Ala Ile Thr Ser Asp Asp Glu Ser Gln Ala Met Gln Thr Val Gly
225 230 235 240

Val His Ser Ile Val Gln Gln Leu His Arg Asn Ser Ile Gln Phe Thr
245 250 255

Asp Gly Tyr Glu Val Lys Glu Asp Ile Gly Val Gly Ser Tyr Ser Val
260 265 270

Cys Lys Arg Cys Ile His Lys Ala Thr Asn Met Glu Phe Ala Val Lys
275 280 285

Val Asn Phe Phe Tyr Leu Lys Cys Asn Ser Tyr Ser Ser Cys Ser Cys
290 295 300

Met Ser Val Pro Val Lys Asn Tyr Thr Pro Leu Val Val Lys Ser Ala
305 310 315 320

Phe Cys Tyr Lys Lys Val Lys Tyr Leu Ala Ser Asp Leu Gln Arg Ser
325 330 335

<210> 175

<211> 198

<212> PRT

<213> Homo sapiens

<400> 175

Met Pro Leu Ala Gln Leu Ala Asp Pro Trp Gln Lys Met Ala Val Glu
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Ser Pro Ser Asp Ser Ala Glu Asn Gly Gln Gln Ile Met Asp Glu Pro
20 25 30

Met Gly Glu Glu Glu Ile Asn Pro Gln Thr Glu Glu Val Ser Ile Lys
35 40 45

Glu Ile Ala Ile Thr His His Val Lys Glu Gly His Glu Lys Ala Asp
50 55 60

Pro Ser Gln Phe Glu Leu Leu Lys Val Leu Gly Gln Gly Ser Phe Gly
65 70 75 80

Lys Val Phe Leu Val Lys Lys Ile Ser Gly Ser Asp Ala Arg Gln Leu
85 90 95

Tyr Ala Met Lys Val Leu Lys Lys Ala Thr Leu Lys Val Arg Asp Arg
100 105 110

Val Arg Thr Lys Met Glu Arg Asp Ile Leu Val Glu Val Asn His Pro
115 120 125

Phe Ile Val Lys Leu His Tyr Ala Phe Gln Thr Glu Gly Lys Leu Tyr
130 135 140

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Leu Ile Leu Asp Phe Leu Arg Gly Gly Asp Leu Phe Thr Arg Leu Ser
145 150 155 160

Lys Glu Val Met Phe Thr Glu Glu Asp Val Lys Phe Tyr Leu Ala Glu
165 170 175

Leu Ala Leu Ala Leu Asp His Leu His Ser Leu Gly Ile Ile Tyr Arg
180 185 190

Asp Leu Lys Pro Glu Lys
195

<210> 176

<211> 489

<212> PRT

<213> Homo sapiens

<400> 176

Met Ser Thr Glu Ala Asp Glu Gly Ile Thr Phe Ser Val Pro Pro Phe
1 5 10 15

Ala Pro Ser Gly Phe Cys Thr Ile Pro Glu Gly Gly Ile Cys Arg Arg
20 25 30

Gly Gly Ala Ala Ala Val Gly Glu Gly Glu Glu His Gln Leu Pro Pro
35 40 45

Pro Pro Pro Gly Ser Phe Trp Asn Val Glu Ser Ala Ala Ala Pro Gly
50 55 60

Ile Gly Cys Pro Ala Ala Thr Ser Ser Ser Ser Ala Thr Arg Gly Arg
65 70 75 80

Gly Ser Ser Val Gly Gly Gly Ser Arg Arg Thr Thr Val Ala Tyr Val
85 90 95

Ile Asn Glu Ala Ser Gln Gly Gln Leu Val Val Ala Glu Ser Glu Ala
100 105 110

Leu Gln Ser Leu Arg Glu Ala Cys Glu Thr Val Gly Ala Thr Leu Glu
115 120 125

Thr Leu His Phe Gly Lys Leu Asp Phe Gly Glu Thr Thr Val Leu Asp
130 135 140

Arg Phe Tyr Asn Ala Asp Ile Ala Val Val Glu Met Ser Asp Ala Phe
145 150 155 160

Arg Gln Pro Ser Leu Phe Tyr His Leu Gly Val Arg Glu Ser Phe Ser
165 170 175

Met Ala Asn Asn Ile Ile Leu Tyr Cys Asp Thr Asn Ser Asp Ser Leu
180 185 190

Gln Ser Leu Lys Glu Ile Ile Cys Gln Lys Asn Thr Met Cys Thr Gly
195 200 205

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Asn	Tyr	Thr	Phe	Val	Pro	Tyr	Met	Ile	Thr	Pro	His	Asn	Lys	Val	Tyr	
210						215				220						
Cys	Cys	Asp	Ser	Ser	Phe	Met	Lys	Gly	Leu	Thr	Glu	Leu	Met	Gln	Pro	
225						230				235					240	
Asn	Phe	Glu	Leu	Leu	Leu	Gly	Pro	Ile	Cys	Leu	Pro	Leu	Val	Asp	Arg	
245						250				255						
Phe	Ile	Gln	Leu	Leu	Lys	Val	Ala	Gln	Ala	Ser	Ser	Ser	Gln	Tyr	Phe	
260						265				270						
Arg	Glu	Ser	Ile	Leu	Asn	Asp	Ile	Arg	Lys	Ala	Arg	Asn	Leu	Tyr	Thr	
275						280				285						
Gly	Lys	Glu	Leu	Ala	Ala	Glu	Leu	Ala	Arg	Ile	Arg	Gln	Arg	Val	Asp	
290						295				300						
Asn	Ile	Glu	Val	Leu	Thr	Ala	Asp	Ile	Val	Ile	Asn	Leu	Leu	Leu	Ser	
305						310				315					320	
Tyr	Arg	Asp	Ile	Gln	Asp	Tyr	Asp	Ser	Ile	Val	Lys	Leu	Val	Glu	Thr	
325						330				335						
Leu	Glu	Lys	Leu	Pro	Thr	Phe	Asp	Leu	Ala	Ser	His	His	His	Val	Lys	
340						345				350						
Phe	His	Tyr	Ala	Phe	Ala	Leu	Asn	Arg	Arg	Asn	Leu	Pro	Gly	Asp	Arg	
355						360				365						
Ala	Lys	Ala	Leu	Asp	Ile	Met	Ile	Pro	Met	Val	Gln	Ser	Glu	Gly	Gln	
370						375				380						
Val	Ala	Ser	Asp	Met	Tyr	Cys	Leu	Val	Gly	Arg	Ile	Tyr	Lys	Asp	Met	
385						390				395					400	
Phe	Leu	Asp	Ser	Asn	Phe	Thr	Asp	Thr	Glu	Ser	Arg	Asp	His	Gly	Ala	
405						410				415						
Ser	Trp	Phe	Lys	Lys	Ala	Phe	Glu	Ser	Glu	Pro	Thr	Leu	Gln	Ser	Gly	
420						425				430						
Ile	Asn	Tyr	Ala	Val	Leu	Leu	Leu	Ala	Ala	Gly	His	Gln	Phe	Glu	Ser	
435						440				445						
Ser	Phe	Glu	Leu	Arg	Lys	Val	Gly	Asn	Tyr	Asn	Leu	Asn	Phe	Tyr	Met	
450						455				460						
Glu	Ile	Lys	Lys	Leu	Gly	Pro	Asn	Leu	Val	Gln	Arg	Arg	Ile	Ser	Ala	
465						470				475					480	
Asp	Ser	Asp	Gly	Ser	Pro	Gly	Phe	Val								
485																

<210> 177

<211> 105

<212> PRT
<213> Homo sapiens

<400> 177

Met	Arg	Glu	Phe	Glu	Val	Leu	Lys	Lys	Leu	Asn	His	Lys	Asn	Ile	Val
1				5					10					15	
Lys	Leu	Phe	Ala	Ile	Glu	Glu	Glu	Thr	Thr	Thr	Arg	His	Lys	Val	Leu
			20					25					30		
Ile	Met	Glu	Phe	Cys	Pro	Cys	Gly	Ser	Leu	Tyr	Thr	Val	Leu	Glu	Glu
		35					40					45			
Pro	Ser	Asn	Ala	Tyr	Gly	Leu	Pro	Glu	Ser	Glu	Phe	Leu	Ile	Val	Leu
	50					55					60				
Arg	Asp	Val	Val	Gly	Gly	Met	Asn	His	Leu	Arg	Glu	Asn	Gly	Ile	Val
65					70					75					80
His	Arg	Asp	Ile	Lys	Pro	Gly	Asn	Ile	Met	Arg	Ala	Leu	Tyr	His	Ser
				85					90					95	
Leu	Val	Asp	Asp	Ser	Phe	His	Pro	Pro							
			100					105							

<210> 178

<211> 413

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(413)

<223> "XAA" can be any amino acid

<400> 178

Met	Tyr	Cys	Phe	Gly	Arg	Lys	Xaa	Tyr	Ile	Ser	Thr	Arg	Pro	Cys	Phe
1				5					10					15	
Pro	Asn	Lys	Thr	Cys	Gln	Lys	Met	Leu	Ile	Ile	Leu	Thr	Ser	Ala	Leu
			20					25					30		
Gln	Ile	Ala	His	Arg	Cys	Ile	Cys	Arg	Ile	Leu	Leu	Gly	Ser	Arg	Val
		35					40					45			
Leu	Ala	Ala	Lys	Ala	Ser	Gly	Asn	Cys	Thr	Leu	Asn	Ser	Glu	Asp	Phe
	50					55				60					
Ile	Phe	Asn	Ile	Gly	Ser	Ala	Ala	Tyr	Asp	Ala	Val	Leu	Asp	Arg	Asn
65					70				75						80
Val	Ala	Ile	Lys	Lys	Leu	Ser	Arg	Pro	Phe	Gln	Asn	Gln	Thr	His	Ala
				85					90					95	

Lys Arg Ala Tyr Arg Glu Leu Val Leu Met Lys Cys Val Asn His Lys
100 105 110

Asn Ile Ile Ser Leu Leu Asn Val Phe Thr Pro Gln Lys Thr Leu Glu
115 120 125

Glu Phe Gln Asp Val Tyr Leu Val Met Glu Leu Met Asp Ala Asn Leu
130 135 140

Cys Gln Val Ile Gln Met Glu Leu Asp His Glu Arg Met Ser Tyr Leu
145 150 155 160

Leu Tyr Gln Met Leu Cys Gly Ile Lys His Leu His Ser Ala Gly Ile
165 170 175

Ile His Arg Asp Leu Lys Pro Ser Asn Ile Val Val Lys Ser Asp Cys
180 185 190

Thr Leu Lys Ile Leu Asp Phe Gly Leu Ala Arg Thr Ala Gly Thr Ser
195 200 205

Phe Met Met Thr Pro Tyr Val Val Thr Arg Tyr Tyr Arg Ala Pro Glu
210 215 220

Val Ile Leu Gly Met Gly Tyr Lys Glu Asn Val Asp Ile Trp Ser Val
225 230 235 240

Gly Cys Ile Met Gly Glu Met Val Arg His Lys Ile Leu Phe Pro Gly
245 250 255

Arg Asp Tyr Ile Asp Gln Trp Asn Lys Val Ile Glu Gln Leu Gly Thr
260 265 270

Pro Cys Pro Glu Phe Met Lys Lys Leu Gln Pro Thr Val Arg Asn Tyr
275 280 285

Val Glu Asn Arg Pro Lys Tyr Ala Gly Leu Thr Phe Pro Lys Leu Phe
290 295 300

Pro Asp Ser Leu Phe Pro Ala Asp Ser Glu His Asn Lys Leu Lys Ala
305 310 315 320

Ser Gln Ala Arg Asp Leu Leu Ser Lys Met Leu Val Ile Asp Pro Ala
325 330 335

Lys Arg Ile Ser Val Asp Asp Ala Leu Gln His Pro Tyr Ile Asn Val
340 345 350

Trp Tyr Asp Pro Ala Glu Val Glu Ala Pro Pro Pro Gln Ile Tyr Asp
355 360 365

Lys Gln Leu Asp Glu Arg Glu His Thr Ile Glu Glu Trp Lys Glu Leu
370 375 380

Ile Tyr Lys Glu Val Met Asn Ser Glu Glu Lys Thr Lys Asn Gly Val
385 390 395 400

Val Lys Gly Gln Pro Ser Pro Ser Ala Gln Val Gln Gln
 405 410

<210> 179
 <211> 108
 <212> PRT
 <213> Homo sapiens

<400> 179

Met Ser Lys Ser Lys Val Asp Asn Gln Phe Tyr Ser Val Glu Val Gly
 1 5 10 15
 Asp Ser Thr Phe Thr Val Leu Lys Arg Tyr Gln Asn Leu Lys Pro Ile
 20 25 30
 Gly Ser Gly Ala Gln Gly Ile Val Cys Ala Ala Tyr Asp Ala Val Leu
 35 40 45
 Asp Arg Asn Val Ala Ile Lys Lys Leu Ser Arg Pro Phe Gln Asn Gln
 50 55 60
 Thr His Ala Lys Arg Ala Tyr Arg Glu Leu Val Leu Met Lys Cys Val
 65 70 75 80
 Asn His Lys Asn Val Ser Phe Val Ile Phe Lys Leu Leu Ala Val Gly
 85 90 95
 Val Cys Lys Ile Gly Lys Arg Lys Cys Val Cys Thr
 100 105

<210> 180
 <211> 336
 <212> PRT
 <213> Homo sapiens

<400> 180

Met Ala Met Thr Gly Ser Thr Pro Cys Ser Ser Met Ser Asn His Thr
 1 5 10 15
 Lys Glu Arg Val Thr Met Thr Lys Val Thr Leu Glu Asn Phe Tyr Ser
 20 25 30
 Asn Leu Ile Ala Gln His Glu Glu Arg Glu Met Arg Gln Lys Lys Leu
 35 40 45
 Glu Lys Val Met Glu Glu Glu Gly Leu Lys Asp Glu Glu Lys Arg Leu
 50 55 60
 Arg Arg Ser Ala His Ala Arg Lys Glu Thr Glu Phe Leu Arg Leu Lys
 65 70 75 80
 Arg Thr Arg Leu Gly Leu Glu Asp Phe Glu Ser Leu Lys Val Ile Gly
 85 90 95
 Arg Gly Ala Phe Gly Glu Val Arg Leu Val Gln Lys Lys Asp Thr Gly

100

105

110

His Val Tyr Ala Met Lys Ile Leu Arg Lys Ala Asp Met Leu Glu Lys
 115 120 125

Glu Gln Val Gly His Ile Arg Ala Glu Arg Asp Ile Leu Val Glu Ala
 130 135 140

Asp Ser Leu Trp Val Val Lys Met Phe Tyr Ser Phe Gln Asp Lys Leu
 145 150 155 160

Asn Leu Tyr Leu Ile Met Glu Phe Leu Pro Gly Gly Asp Met Met Thr
 165 170 175

Leu Leu Met Lys Lys Asp Thr Leu Thr Glu Glu Glu Thr Gln Phe Tyr
 180 185 190

Ile Ala Glu Thr Val Leu Ala Ile Asp Ser Ile His Gln Leu Gly Phe
 195 200 205

Ile His Arg Asp Ile Lys Pro Asp Asn Leu Leu Leu Asp Ser Lys Gly
 210 215 220

His Val Lys Leu Ser Asp Phe Gly Leu Cys Thr Gly Leu Lys Lys Ala
 225 230 235 240

His Arg Thr Glu Phe Tyr Arg Asn Leu Asn His Ser Leu Pro Ser Asp
 245 250 255

Phe Thr Phe Gln Asn Met Asn Ser Lys Arg Lys Ala Glu Thr Trp Lys
 260 265 270

Arg Asn Arg Arg Gln Leu Ala Phe Ser Thr Val Gly Thr Pro Asp Tyr
 275 280 285

Ile Ala Pro Glu Val Phe Met Gln Thr Gly Tyr Asn Lys Leu Cys Asp
 290 295 300

Trp Trp Ser Leu Gly Val Ile Met Tyr Glu Met Leu Ile Gly Lys Leu
 305 310 315 320

His Gly Phe Arg Gly Leu Phe Leu Cys Ile His Asp Arg Leu Leu His
 325 330 335

<210> 181

<211> 415

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(415)

<223> "XAA " can be any amino acid

<400> 181

Xaa Arg His Glu Ser Ala Arg Ala Ala Arg Val Ser Gly Gly Ser Met
1 5 10 15

Leu Asp Ile Ile Lys Tyr Ile Val Asn Arg Gly Glu His Lys Asn Gly
20 25 30

Val Leu Glu Glu Ala Ile Ile Ala Thr Ile Leu Lys Glu Val Leu Glu
35 40 45

Gly Leu Asp Tyr Leu His Arg Asn Gly Gln Ile His Arg Asp Leu Lys
50 55 60

Ala Gly Asn Ile Leu Leu Gly Glu Asp Gly Ser Val Gln Ile Ala Asp
65 70 75 80

Phe Gly Val Ser Ala Phe Leu Ala Thr Gly Gly Asp Val Thr Arg Asn
85 90 95

Lys Val Arg Lys Thr Phe Val Gly Thr Pro Cys Trp Met Ala Pro Glu
100 105 110

Val Met Glu Gln Val Arg Gly Tyr Asp Phe Lys Ala Asp Met Trp Ser
115 120 125

Phe Gly Ile Thr Ala Ile Glu Leu Ala Thr Gly Ala Ala Pro Tyr His
130 135 140

Lys Tyr Pro Pro Met Lys Val Leu Met Leu Thr Leu Gln Asn Asp Pro
145 150 155 160

Pro Thr Leu Glu Thr Gly Val Glu Asp Lys Glu Met Met Lys Lys Tyr
165 170 175

Gly Lys Ser Phe Arg Lys Leu Leu Ser Leu Cys Leu Gln Lys Asp Pro
180 185 190

Ser Lys Arg Pro Thr Ala Ala Glu Leu Leu Lys Cys Lys Phe Phe Gln
195 200 205

Lys Ala Lys Asn Arg Glu Tyr Leu Ile Glu Lys Leu Leu Thr Arg Thr
210 215 220

Pro Asp Ile Ala Gln Arg Ala Lys Lys Val Arg Arg Val Pro Gly Ser
225 230 235 240

Ser Gly His Leu His Lys Thr Glu Asp Gly Asp Trp Glu Trp Ser Asp
245 250 255

Asp Glu Met Asp Glu Lys Ser Glu Glu Gly Lys Ala Ala Phe Ser Gln
260 265 270

Glu Lys Ser Arg Arg Val Lys Glu Glu Asn Pro Glu Ile Ala Val Ser
275 280 285

Ala Ser Thr Ile Pro Glu Gln Ile Gln Ser Leu Ser Val His Asp Ser
290 295 300

Gln Gly Pro Pro Asn Ala Asn Glu Asp Tyr Arg Glu Ala Ser Ser Cys
305 310 315 320

Ala Val Asn Leu Val Leu Arg Leu Arg Asn Ser Arg Lys Glu Leu Asn
325 330 335

Asp Ile Arg Phe Glu Phe Thr Pro Gly Arg Asp Thr Ala Asp Gly Val
340 345 350

Ser Gln Glu Leu Phe Ser Ala Gly Leu Val Asp Gly His Asp Val Val
355 360 365

Ile Val Ala Ala Asn Leu Gln Lys Ile Val Asp Asp Pro Lys Ala Leu
370 375 380

Lys Thr Leu Thr Phe Lys Leu Ala Ser Gly Cys Asp Gly Ser Glu Ile
385 390 395 400

Pro Asp Glu Val Lys Leu Ile Gly Phe Ala Gln Leu Ser Val Ser
405 410 415

<210> 182
<211> 409
<212> PRT
<213> Homo sapiens

<220>
<221> -
<222> (1)..(409)
<223> "Xaa" can be any amino acid

<400> 182

Xaa Arg His Glu Ser Ala Arg Ala Ala Arg Val Ser Gly Gly Ser Met
1 5 10 15

Leu Asp Ile Ile Lys Tyr Ile Val Asn Arg Gly Glu His Lys Asn Gly
20 25 30

Val Leu Glu Glu Ala Ile Ile Ala Thr Ile Leu Lys Glu Val Leu Glu
35 40 45

Gly Leu Asp Tyr Leu His Arg Asn Gly Gln Ile His Arg Asp Leu Lys
50 55 60

Ala Gly Asn Ile Leu Leu Gly Glu Asp Gly Ser Val Gln Ile Ala Asp
65 70 75 80

Phe Gly Val Ser Ala Phe Leu Ala Thr Gly Gly Asp Val Thr Arg Asn
85 90 95

Lys Val Arg Lys Thr Phe Val Gly Thr Pro Cys Trp Met Ala Pro Glu
100 105 110

Val Met Glu Gln Val Arg Gly Tyr Asp Phe Lys Ala Asp Met Trp Ser
115 120 125

